



# STIC Search Report

## Biotech-Chem Library

STIC Database Tracking Number: 174625

TO: Phillip Gambel  
Location: REM/3E81/3C70  
Art Unit: 1644 22  
Friday, December 2, 2005  
Case Serial Number: 09/751797

From: Toby Port  
Location: Biotech-Chem Library  
REM-1A59  
Phone: 571-272-2523  
[toby.port@uspto.gov](mailto:toby.port@uspto.gov)

### Search Notes

Examiner Gambel,

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Toby Port  
X22523

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From: Chan, Christina  
Sent: Monday, December 19, 2005 3:54 PM  
To: Gambel, Phillip; STIC-Biotech/ChemLib  
Subject: RE: 09751797 interference search PLEASE RUSH SEQUENCE INTERFERENCE SEARCH

Please rush. Thanks Chris

Chris Chan  
TC 1600 New Hire Training Coordinator and SPE 1644  
(571)-272-0841  
Remsen, 3E89

-----Original Message-----

From: Gambel, Phillip  
Sent: Monday, December 19, 2005 3:44 PM  
To: Chan, Christina  
Subject: 09751797 interference search PLEASE RUSH SEQUENCE INTERFERENCE SEARCH

christina

PLEASE RUSH SEQUENCE INTERFERENCE SEARCH for **09 / 751,797** AFTER FINAL

TO STIC-Biotech/ChemLib

please perform a **SEQUENCE INTERFERENCE SEARCH** for **09 / 751,797** in PAPER

against PENDING and U.S. Patents, PGPubs

SEQ ID NO: 7

SEQ ID NO: 8

SEQ ID NO 9

SeQ ID NO: 24

SEQ ID NO: 25

SEQ ID NO: 29

RECEIVED  
DEC 19 2005  
STIC/STIC-INT-2005

\*\*\*\*\*

Searcher: \_\_\_\_\_  
Searcher Phone: \_\_\_\_\_  
Date Searcher Picked up: \_\_\_\_\_  
Date completed: \_\_\_\_\_  
Searcher Prep Time: \_\_\_\_\_  
Online Time: \_\_\_\_\_

\*\*\*\*\*

Type of Search  
NA# \_\_\_\_\_ AA# \_\_\_\_\_  
S/L: \_\_\_\_\_ Oligomer: \_\_\_\_\_  
Encode/Transl: \_\_\_\_\_  
Structure #: \_\_\_\_\_ Text: \_\_\_\_\_  
Inventor: \_\_\_\_\_ Litigation: \_\_\_\_\_

\*\*\*\*\*

Vendors and cost where applicable  
STN: \_\_\_\_\_  
DIALOG: \_\_\_\_\_  
QUESTEL/ORBIT: \_\_\_\_\_  
LEXIS/NEXIS: \_\_\_\_\_  
SEQUENCE SYSTEM: \_\_\_\_\_  
WWW/Internet: \_\_\_\_\_  
Other (Specify): \_\_\_\_\_

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GenCore version 5.1.6  
Copyright (c) 1993 - 2005 CompuGen Ltd.

OM/nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 126.661 Seconds  
(without alignments)  
15704.028 Million cell updates/sec

Title: US-09-751-797-7  
Perfect score: 1119  
Sequence: 1 taacagggtctctctctcac.....tggatcataaaaaaaa 1119

Scoring table: IDENTITY NUC  
Gapop 10.0 , Gapext 1.0

Searched: 1303057 seqs, 888780828 residues

Total number of hits satisfying chosen parameters: 2606114

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Issued Patents NA: \*  
1: /cgn2\_6/ptodata/1/ina/1\_COMB.seq: \*  
2: /cgn2\_6/ptodata/1/ina/5\_COMB.seq: \*  
3: /cgn2\_6/ptodata/1/ina/6A\_COMB.seq: \*  
4: /cgn2\_6/ptodata/1/ina/6B\_COMB.seq: \*  
5: /cgn2\_6/ptodata/1/ina/H\_COMB.seq: \*  
6: /cgn2\_6/ptodata/1/ina/PCUS\_COMB.seq: \*  
7: /cgn2\_6/ptodata/1/ina/PP\_COMB.seq: \*  
8: /cgn2\_6/ptodata/1/ina/RE\_COMB.seq: \*  
9: /cgn2\_6/ptodata/1/ina/backfiles1.seq: \*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1119	100.0	1119	3	US-09-178-973B-7
2	1119	100.0	1119	3	US-09-419-568F-7
3	1119	100.0	1119	3	US-09-354-243B-7
4	1107.8	99.0	1166	3	US-10-084-298-3
5	1047.8	93.6	1111	3	US-09-178-973B-9
6	1047.8	93.6	1111	3	US-09-419-568F-9
7	1047.8	93.6	1111	3	US-09-354-243B-9
8	993.2	88.8	1050	3	US-10-090-365-40
9	601.4	53.7	7445	3	US-09-178-973B-8
10	601.4	53.7	7445	3	US-09-419-568F-8
11	601.4	53.7	7445	3	US-09-354-243B-8
12	555.2	49.6	5935	3	US-09-178-973B-17
13	555.2	49.6	5935	3	US-09-419-568F-29
14	555.2	49.6	5935	3	US-09-354-243B-29
15	535.2	47.8	1191	3	US-10-084-298-1
16	524.8	46.9	1152	3	US-09-870-574-1
17	524.4	46.9	1116	3	US-10-090-365-14
18	524.4	46.9	1116	3	US-09-728-911-14
19	409.2	36.6	689	3	US-09-949-016-5443
20	409.2	36.6	690	3	US-09-419-568F-24
21	409.2	36.6	690	3	US-09-354-243B-24
22	216.6	19.4	8888	3	US-09-949-016-17185
23	186.2	16.6	191	3	US-10-084-298-9
24	126	11.3	4797	3	US-09-419-568F-25

ALIGNMENTS

RESULT 1

US-09-178-973B-7

; Sequence 7, Application US/09178973B

; Patent No. 6274710

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Louhed, Jamila

; APPLICANT: Renauld, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; TITLE OF INVENTION: (TIFFS)

; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5543

; CURRENT APPLICATION NUMBER: US/09/178,973B

; CURRENT FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 17

; SEQ ID NO 7

; LENGTH: 1119

; TYPE: DNA

; ORGANISM: Mus musculus

US-09-178-973B-7

Query Match 100.0%; Score 1119; DB 3; Length 1119;

Best Local Similarity 100.0%; Pred. No. 3.3e-272;

Matches 1119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy	61	CTGCAGAAATCTATGAGTTTTCCTTATGGGACTTTGGCCGACGTCGCTGCTCTC	120
Db	61	CTGCAGAAATCTATGAGTTTTCCTTATGGGACTTTGGCCGACGTCGCTGCTCTC	120
Qy	121	ATTGCCCTGTGGCCGACGAGGCAATGCGTCGCGTCAACACCCGCGTGAAGCTTGAG	180
Db	121	ATTGCCCTGTGGCCGACGAGGCAATGCGTCGCGTCAACACCCGCGTGAAGCTTGAG	180
Qy	181	GTGTCCTCACTTCCAGCAGCGGTACATGTCGTCGCGTCAACACCCGCGTGAAGCTTGAG	240
Db	181	GTGTCCTCACTTCCAGCAGCGGTACATGTCGTCGCGTCAACACCCGCGTGAAGCTTGAG	240
Qy	241	AGCTTTCAGATAAACAACAGACGTCGCGCTCATCGGGAGAGAACTGTTCCGAGGAGTC	300
Db	241	AGCTTTCAGATAAACAACAGACGTCGCGCTCATCGGGAGAGAACTGTTCCGAGGAGTC	300
Qy	301	AGTGCTAAAGATCAGTGCTACCTGATGAACAGGTCCTCACTTACCTTCCGAGGAGCTT	360
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Db 361 CTGCTCCCGCAGTCAGACAGGTTCCAGCCCTACATGACAGAGGTGGTACCTTTCTTGACC 420  
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Db 421 AAACCTCAGCAATCAGCTCAGCTCCTGTGCATCATCAGCGGTGACGACCAAGACATCCAGAAG 480  
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Db 541 ATTGGGAACTGGACCTGCTGTTTATGTCTGTGAGAAATGCTTGGCTGTGAGCGAGAGA 600  
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Db 661 TGGACTTTTTTACTTAAAGGAAAGTGAAGGCTAACGCTCCATCATATTAGAAGATTTTAC 720  
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Db 721 ATGAAACCTGGCTCAGTTGAAAGAAATAGTGTCAAGTTGCTCCATGAGACCCAGAGTA 780  
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Qy 841 AAATAATGTACTTTTAAAGAAATTTTGAAGAGGTTTACCTCTCATTTCTTTAGAAAAA 900  
Db 841 AAATAATGTACTTTTAAAGAAATTTTGAAGAGGTTTACCTCTCATTTCTTTAGAAAAA 900  
Qy 901 AGCTTATGTAACCTTCCATTCATCCAAATATTTTATTTATGTAAGTTTATTTATATA 960  
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Qy 1081 TTAGGCTTTAATAACACATGGATATCATATAAAAAA 1119  
Db 1081 TTAGGCTTTAATAACACATGGATATCATATAAAAAA 1119

RESULT 2  
US-09-419-568F-7  
; Sequence 7, Application US/09419568F  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/419,568F  
; CURRENT FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29

; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-419-568F-7  
  
Query Match 100.0%; Score 1119; DB 3; Length 1119;  
Best Local Similarity 100.0%; Pred. No. 3.3e-272;  
Matches 1119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
Qy 1 TAAACAGGCT 60  
Db 1 TAAACAGGCT 60  
Qy 61 CTGAGAGAAATCTATGAGTGTCT 120  
Db 61 CTGAGAGAAATCTATGAGTGTCT 120  
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Qy 181 GTGTCCAACTTCCAGCAGCGGTACATCGTCAACCGCACTTTTATGTGGCCAAAGAGGCC 240  
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Qy 241 AGCTTTGAGATTAACACAGACAGCTCGGCTCATCGGGAGAAACTGTTCCGAGAGTC 300  
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Db 361 CTGCTCCCGCAGTCAGACAGGTTCCAGCCCTACATGACAGAGGTGGTACCTTTCTTGACC 420  
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Qy 601 AGCTAGAAACGGAAGAACTGCT 660  
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Db 661 TGGACTTTTTTACTTAAAGGAAAGTGAAGGCTAACGCTCCATCATATTAGAAGATTTTAC 720  
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Db 781 GACTTGATAACCAAGATTCATTGCAATATTTTATTTTATTTGCTGATGATACAAAGAA 840  
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Db 841 AAATAATGTACTTTTAAAGAAATTTTGAAGAGGTTTACCTCTCATTTCTTTAGAAAAA 900  
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Db 961 GTATACATTTTATTTATGTCAGTTTATTAATATGAGTTTATTTATAGAAAACATTATCTGC 1020  
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Db 1021 TATGATATTTAGTATAGGCAATATATATTTATGACATTAACATGGAACCAAGATATC 1080  
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RESULT 3  
US-09-354-243B-7  
; Sequence 7, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Pa  
; TITLE OF INVENTION: (Tifs)  
; FILE REFERENCE: LUD 5543.1  
; CURRENT FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-354-243B-7

Query Match 100.0%; Score 1119; DB 3; Length 1119;  
Best Local Similarity 100.0%; Pred. No. 3.3e-272;  
Matches 1119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1081 TTAGGCTTTAATAAACAATGATATCATATAAAAAA 1119

RESULT 4  
US-10-084-298-3  
; Sequence 3, Application US/10084298  
; Patent No. 6939545  
; GENERAL INFORMATION:  
; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: Xuan, Dejun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory  
; TITLE OF INVENTION: Disorders  
; FILE REFERENCE: G15358 CIP  
; CURRENT APPLICATION NUMBER: US/10/084,298  
; CURRENT FILING DATE: 2002-09-10  
; PRIOR APPLICATION NUMBER: 60/270,823  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR APPLICATION NUMBER: 60/281,353  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR APPLICATION NUMBER: 60/131,473  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR APPLICATION NUMBER: 09/561,811  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 3

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; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Murine
US-10-084-298-3

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Query Match 99.0%; Score 1107.8; DB 3; Length 1166;  
Best Local Similarity 99.4%; Pred. No. 2.2e-269;  
Matches 1112: Conservative 0; Mismatches 7; Indels 0;

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84	Db				143
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144	Db				203
181	Qy	GTG	TCAA	CTTCAGACGCCGTACATCGTCAACCGCACCTTTATGCTGGCCGCAAGAGGCC	240
204	Db				263
241	Qy	AGC	CTGCAGATA	CAACACACAGAGCTCGCGCTCATCGGGAGAAACTGTTCCGAGGAGTC	300
264	Db				323
301	Qy	AGT	GCTAAAGAT	CACTGATGAAGCAGGTGCTCAACTTCACCCCTGGAAGACGTT	360
324	Db				383
361	Qy	CTG	TCCCCCAGT	CAGACAGGTTCCAGCCCTACATGACAGGAGTGGTACCTTTCCTGACC	420
384	Db				443
421	Qy	AAA	CTCAGCANT	CAGCTCAGCTCCGTCCATCATCAGCGGTGACGACAGAACATCCAGAG	480
444	Db				503
481	Qy	AAT	GTCAGAAGC	CTGAAGGACAGTGA AAAAGCTTTGGAGAGAGTGGAGAGATCAAGGCG	540
504	Db				563
541	Qy	ATT	GGGAACT	TGGACCTGCTGTTTATGTCCTGAGAAATGCTTGCGCTGACGCGAGAAGA	600
564	Db				623
601	Qy	AGC	TAGAAA	CGAAGAACTGCTCCTTCTGCTCCCTTTCTA AAAAGAA CAATAAGATCCCTGAA	660
624	Db				683
661	Qy	TGG	ACTTTTTT	TACTAAAGAAAGTCAGAGCTTAAAGCTCCATCATCATTTAGAAAGATTTCCAC	720
684	Db				743
721	Qy	ATG	AAACCT	TGCGCTCAGTTGAAAAGAAATAGTGTCAAAGTTGTCATGAGACAGAGGTA	780
744	Db				803
781	Qy	GAC	TTGATA	CCACCAAGATTCATTTGACAAATTTTTATTTGTCACCTGATGATCAACAGAA	840
804	Db				863
841	Qy	AAA	TAACTG	ACTTTTTAAAAAATGGTTTGAAGAGAGGTTTACCTCTCATCTCCTTTAGAAAAA	900
864	Db				923
901	Qy	AGC	TTATG	TAACTTCATTTCCATATCCAAATATTTTATATATGATGAAGTTATTTATPATAA	960
924	Db				983

## RESULT 5

US-09-178-973B-9  
: Sequence 9, Application US/09178973B

; Patent No. 6274710

; GENERAL INFORMATION:

**APPLICANT: Dumoutier, Laure**

APPLICANT: Louhed, Jamila

APPLICANT: Renault, Jean-Christophe

**TITLE OF INVENTION: Isolate**

1. TITLE OF INVENTION: (TIFS)

**TITLE OF INVENTION:** The Proteins Encoded,

FILE REFERENCE: LUD 5543

: CURRENT APPLICATION NUMBER

CONCURRENT

; CURRENT PLANNING  
; NUMBER OF SECTIONS; NUMBER OF  
: CEO ID NO 9

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; SEQ ID NO 9
      LENGTH: 1111

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; LENGTH: 111
; TYPE: DNA

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Overall Match 03 68: score 1047 8: DB 3: length 1111:

Query Match

Post Local Similarity	97.08	Pred	No	39-254	DE
		Score	1047.8		

BEST LOCAL SIMILARITY 97.0%; PRED. NO: 3E-234;  
MATCHES 1078. CONSENSUATIVE 0. MISMATCHES 32. INDEX 1. GAPS 1.

QY	3	AACAGGCTCTCCTCTCACTTAACTGTGACACTTGTGCGATCTGTAGTGGCTGTGCTCT	62
DB	1	AACAGGCTCTCCTCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGGCTGTGCTCT	60
QY	63	GCAGAAATCTATGAGTTTTTCCCCTTATGGGGACATTTGGCCGACGTGCGTCTCTTCTCAT	122
DB	61	GCAGAAATCTATGAGTTTTTCCCCTTATGGGGACATTTGGCCGACGTGCGTCTCTTCTCAT	120
QY	123	TGCCCTGTGGGCCCAGGAGGCAAAATCGCTCTCCCGTCAACACCCCGGTGCAAGCTTGAGGT	182
DB	121	TGCCCTGTGGGCCCAGGAGGCAAAATCGCTCTCCCGTCAACACCCCGGTGCAAGCTTGAGGT	180
QY	183	GTCCAACTTTCAGCAGCGGTACATCGTCAAACCGCACCTTTATGCTGGCCCAAGAGGCCAG	242
DB	181	GTCCAACTTTCAGCAGCGCGTACATCGTCAAACCGCACCTTTATGCTGGCCCAAGAGGCCAG	240
QY	243	CTTTGCAGATAACAAACAGACGTCGCGCTCATCGGGAGAAACTGTTCCGAGGAGTCAG	302
DB	241	CTTTGCAGATAACAAACAGACGTCGCGCTCATCGGGAGAAACTGTTCCGAGGAGTCAG	300
QY	303	TGCTTAAGATCAGTGCTACCTGATGAACAGGTGCTCAACTTCACCTCGGAGAGCGTTCT	362
DB	301	TGCTTAAGATCAGTGCTACCTGATGAACAGGTGCTCAACTTCACCTCGGAGAGCAATCT	360
QY	363	GCTCCCCCAGTCAGACAGGTTCCAGCCCTACATGACGAGGAGTGATACCTTTCTTGACCAA	422
DB	361	GCTCCCCCAGTCAGACAGGTTCCGSCCTTACATGACGAGGAGTGATGCTTTCTTGACCAA	420
QY	423	ACTCAGCAATCAGCTCAGCTCTGTTCATCATCAGCCGTGACGACCCAGACATCCAGAGAA	482
DB	421	ACTCAGCAATCAGCTCAGCTCTGTTCATCATCAGTGGTGAACACCCAGAACATCCAGAGAA	480
QY	483	TGTCAGAAAGGCTGAAGGAGACAGTCAAAAAGCTTGGAGAGAGTGGAGAGATCAAGGCGAT	542
DB	481	TGTCAGAAAGGCTGAAGGAGACAGTCAAAAAGCTTGGAGAGAGTGGAGAGATCAAGGCGAT	540



RESULT 7

US-09-354-243B-9  
; Sequence 9, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Peptides  
; TITLE OF INVENTION: (TIFFs)  
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.1  
; CURRENT APPLICATION NUMBER: US/09/354,243B  
; CURRENT FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
; US-09-354-243B-9

Query Match 93.6%; Score 1047.8; DB 3; Length 1111;  
Best Local Similarity 97.0%; Pred. No. 3e-254;  
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;

Qy	3	AACAGGCTCTCCTCTCACTTATCAACTGTGTGACACTTGTGCGATCTCTGATGCTGTCCT	62
Db	1	AACAGGCTCTCCTCTCACTTATCAACTGTGTGACACTTGTGCGATCTCTGATGCTGTCCT	60
Qy	63	GCAGAAATCTATGAGTTTTTCCCTTATGGGACCTTTGGCCGCCAGCTGCTGCTCTCAT	122
Db	61	GCAGAAATCTATGAGTTTTTCCCTTATGGGACCTTTGGCCGCCAGCTGCTGCTCTCAT	120
Qy	123	TGCCCTGTGGGCCCAGGAGGCAATGCGCTGCCCGTCAACACCCGGTGCAGGCTTGAGGT	182
Db	121	TGCCCTGTGGGCCCAGGAGGCAATGCGCTGCCCGTCAACACCCGGTGCAGGCTTGAGGT	180
Qy	183	GTCCAACTTCCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGCGCAGAGGCCAG	242
Db	181	GTCCAACTTCCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGCGCAGAGGCCAG	240
Qy	243	CCTTGCAGATAACAAACAGACGTCCTCGGCTCATCGGGAGAACTGTTCCGAGAGTCAG	302
Db	241	CCTTGCAGATAACAAACAGACGTCCTCGGCTCATCGGGAGAACTGTTCCGAGAGTCAG	300
Qy	303	TGCTAAAGATCAGTGCTACCTGTATGAAGCAGGTGCTCAACTTCACTTCCGAAAGCGTTCT	362
Db	301	TGCTAAAGATCAGTGCTACCTGTATGAAGCAGGTGCTCAACTTCACTTCCGAAAGCATTTCT	360
Qy	363	GCTCCCCAGTCAGACAGGTTCCAGCCCTATATGCGAGAGGTGCTACCTTCTGACCAA	422
Db	361	GCTCCCCAGTCAGACAGGTTCCAGCCCTATATGCGAGAGGTGCTACCTTCTGACCAA	420
Qy	423	ACTCAGCAATCAGCTCAGCTCTCTGCATCAGCGGTGACGACCCAGAACATCCAGAAGAA	482
Db	421	ACTCAGCAATCAGCTCAGCTCTCTGCATCAGTGTTGACGACCCAGAACATCCAGAAGAA	480
Qy	483	TGT CAGAAGGCTGAAGGAGACAGTGAAAAAGCTTTGGAGAGAGTGGAAGATCAAGGCGAT	542
Db	481	TGT CAGAAGGCTGAAGGAGACAGTGAAAAAGCTTTGGAGAGAGTGGAAGATCAAAAGCGAT	540
Qy	543	TGGGGAACTGCACTGCTGTTTATGCTCTCAGAGAAATGCTTGGCTCTGCGGAGAGAAAG	602
Db	541	CGGGGAACTGCACTGCTGTTTATGCTCTCAGAGAAATGCTTGGCTCTGCGGAGAGAAAG	600
Qy	603	CTAGAAAACGAAGAACTGCTCTCTTCCCTGCTCTTAAAAAGAACAAATAGATCCCTGAATG	662
Db	601	CTAGAAAACGAAGAACTGCTCTCTTCCCTGCTCTTAAAAAGAACAAATAGATCCCTGAATG	660

123 TGCCCTGTGGCCCGCAGAGGCAAAATGCGTCCCGTCAACACCCGGTGAAGCTTGAGGT 182  
Db TGCCCTGTGGCCCGCAGAGGCAAAATGCGTCCCGTCAACACCCGGTGAAGCTTGAGGT 180  
Qy GTCCAACTTCAGAGCCGCTACATGCTCAACCGCACCTTTATGCTGGCCAGAGGCCAG 242  
Db GTCCAACTTCAGAGCCGCTACATGCTCAACCGCACCTTTATGCTGGCCAGAGGCCAG 240  
Qy CCTTGAGATAAACAACAGACAGCTCCGGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 302  
Db CCTTGAGATAAACAACAGACAGCTCCGGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 300  
Qy TGCTAAGATCAGTGTCTACCTGATGAAGAGCGTCTCAACTTCACTCCCTGGAAGACGTTCT 362  
Db TGCTAAGATCAGTGTCTACCTGATGAAGAGCGTCTCAACTTCACTCCCTGGAAGACATCT 360  
Qy GCTCCCGCCAGTCAGAGGTTCCAGCCCTACATGCGAGGAGTGGTCTTCTGACCAA 422  
Db GCTCCCGCCAGTCAGAGGTTCCAGCCCTACATGCGAGGAGTGGTCTTCTGACCAA 420  
Qy ACTCAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGACGACCCAGAAATCCAGAGAA 482  
Db ACTCAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGACGACCCAGAAATCCAGAGAA 480  
Qy TGTCAGAGGCTGAAGAGACAGTGAAGAAAGCTTGGAGAGAGTGGAGAGATCAAGCGAT 542  
Db TGTCAGAGGCTGAAGAGACAGTGAAGAAAGCTTGGAGAGAGTGGAGAGATCAAGCGAT 540  
Qy TGGGAACTGACCTGCTGTTTATGCTCTGAGAAATGCTTGGTCTGAGCGAGAGAG 602  
Db TGGGAACTGACCTGCTGTTTATGCTCTGAGAAATGCTTGGTCTGAGCGAGAGAG 600  
Qy CTAGAAAACGAAGACTGCTCCTTCTGCTGCTCTTAAAGAAACAATAAGATCCCTGAATG 662  
Db CTAGAAAACGAAGACTGCTCCTTCTGCTGCTCTTAAAGAAACAATAAGATCCCTGAATG 660  
Qy GACTTTTTTCTAAGAGAAAGTGAGAGCTAACGCTCCATCATTTAGAAAGATTTACAT 722  
Db GACTTTTTTCTAAGAGAAAGTGAGAGCTAACGCTCCATCATTTAGAAAGATTTACAT 720  
Qy GAAACCTGGCTCAGTTGAAAAGAAATAGTGTCAAGTTCCTGACGACGAGGTAGA 782  
Db GAAACCTGGCTCAGTTGAAAAGAAATAGTGTCAAGTTCCTGACGACGAGGTAGA 780  
Qy CTGTGATAACCAAGAGATTCATTGACAAATTTTATTTGTCATGATTAATGCAACAGAAA 842  
Db CTGTGATAACCAAGAGATTCATTGACAAATTTTATTTGTCATGATTAATGCAACAGAAA 840  
Qy ATAATGTAATTTTAAAGAAATTTGTTGAAAGGAGGTACCTCTCATTTTGAAGAAAAG 902  
Db ATAATGTAATTTTAAAGAAATTTGTTGAAAGGAGGTACCTCTCATTTTGAAGAAAAG 900  
Qy CTATGTAATCTCATTTCCATATCCAAATATTTATATATGTAAGTTTATTTATTAAGT 962  
Db CTATGTAATCTCATTTCCATATCCAAATATTTATATATGTAAGTTTATTTATTAAGT 960  
Qy ATACATTTTATTTATGTCAGTTTATTAATATGATTTTATTTATAGAAACATTTCTGCTA 1022  
Db ATACATTTTATTTATGTCAGTTTATTAATATGATTTTATTTATAGAAACATTTCTGATG 1020  
1023 TTGATATTT-AGTATAGGCAATAATATT 1051  
Db TTGATATTTGATATAGGCAATAATATT 1050

RESULT 9  
US-09-178-973B-8  
; Sequence 8, Application US/09178973B  
; Patent No. 6274710  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE REFERENCE: LUD 5543  
; CURRENT APPLICATION NUMBER: US/09/178,973B  
; CURRENT FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 17  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
US-09-178-973B-8

Query Match 53.7%; Score 601.4; DB 3; Length 7445;  
Best Local Similarity 99.8%; Pred. No. 2.6e-141;  
Matches 602; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
Qy 510 AAAGCTTGGAGAGAGTGGAGAGATCAAGGCGATGGGAACTGGAACTGCTGTTATGTC 569  
Db 6535 ATAGCTTGGAGAGAGTGGAGAGATCAAGGCGATGGGAACTGGAACTGCTGTTATGTC 6594  
Qy 570 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAAGAACTGCTCTTCT 629  
Db 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAAGAACTGCTCTTCT 6654  
Qy 630 GCCTTCTTAAAGAAACAATAAGATCCCTGAATGGACTTTTTTACTAAAGGAAAGTGAGAA 689  
Db 6655 GCCTTCTTAAAGAAACAATAAGATCCCTGAATGGACTTTTTTACTAAAGGAAAGTGAGAA 6714  
Qy 690 GCTAAGCTCCATCATATTAGAAAGATTTCAATGAAACCTGGCTCAGTTGAAAGAAAA 749  
Db 6715 GCTAAGCTCCATCATATTAGAAAGATTTCAATGAAACCTGGCTCAGTTGAAAGAAAA 6774  
Qy 750 TAGTGTCAAGTTCTCCATGAGACGAGAGTAGACTTGATAACCAAGAGATTCATTGACA 809  
Db 6775 TAGTGTCAAGTTCTCCATGAGACGAGAGTAGACTTGATAACCAAGAGATTCATTGACA 6834  
Qy 810 ATATTTTATTTGTCACATGATGATACACAGAAAAAATAATGTACTTTTAAAAAATTTGTTGAA 869  
Db 6835 ATATTTTATTTGTCACATGATGATACACAGAAAAAATAATGTACTTTTAAAAAATTTGTTGAA 6894  
Qy 870 AGAGGTTACCTCTCATTTCTTTAGAAAAAAGCTTATGTAACCTTTCCATATCCAA 929  
Db 6895 AGAGGTTACCTCTCATTTCTTTAGAAAAAAGCTTATGTAACCTTTCCATATCCAA 6954  
Qy 930 TATTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 989  
Db 6955 TATTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 7014  
Qy 990 ATATGGAATTTATTTATAGAAACATTTATCTGCTATTTGATATTTTAGTATAAGGCAATAATA 1049  
Db 7015 ATATGGAATTTATTTATAGAAACATTTATCTGCTATTTGATATTTTAGTATAAGGCAATAATA 7074  
Qy 1050 TTTATGACAAATACTATGGAACAAGATATCTTAGGCTTTTAAACACATGGAATATCAT 1109  
Db 7075 TTTATGACAAATACTATGGAACAAGATATCTTAGGCTTTTAAACACATGGAATATCAT 7134  
Qy 1110 AAA 1112  
Db 7135 AAA 7137

RESULT 10  
US-09-419-568F-8  
; Sequence 8, Application US/09419568F  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFS) The Proteins Encoded, and Uses Thereof







; TITLE OF INVENTION: (TIPS)  
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543  
; CURRENT APPLICATION NUMBER: US/09/178,973B  
; CURRENT FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 17  
; SEQ ID NO 17  
; LENGTH: 5935  
; TYPE: DNA  
; ORGANISM: Mus musculus  
US-09-178-973B-17

Query Match 49.6%; Score 555.2; DB 3; Length 5935;  
Best Local Similarity 96.0%; Pred. No. 1.1e-129;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
  
QY 510 AAAGCTTGGAGAGAGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 569  
DB 5221 ATAGCTTGGAGAGAGCGGAGAGATCAAGCGATCGGGGAACCTGGACCTGCTGTTTATGTC 5280  
  
QY 570 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAAACGAAAGAACTGCTCCTTCCT 629  
DB 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAAACGAAAGAACTGCTCCTTCCT 5340  
  
QY 630 GCCTTCTAAAAGAACAAATAGATCCCTGATGCACTTTTCTAAGGAAAAGTGAGAA 689  
DB 5341 GCCTTCTAAAAGAACAAATAGATCCCTGATGCACTTTTCTAAGGAAAAGTGAGAA 5400  
  
QY 690 GCTAACCTGCCATCATCATTTAGAGATTTTCATGAAACCTGGCTCAGTTGAAAAGAGAAA 749  
DB 5401 GCTAACCTGCCATCATTTAGAGATTTTCATGAAACCTGGCTCAGTTGAAAAGAGAAA 5460  
  
QY 750 TAGTGTCACAGTTGTCATGAGACCCAGAGGTAGACTTGATTAACCAACAAAGATTTCATTGACA 809  
DB 5461 TAGTGTCACAGTTGTCATGAGACCCAGAGGTAGACTTGATTAACCAACAAAGATTTCATTGACA 5520  
  
QY 810 ATATTTTATGTCAGTATGATACACAGAAAATATATGTAATCTTAAAAAATGTTTGAA 869  
DB 5521 ATATTTTATGTCAGTATGATACACAGAAAATATATGTAATCTTAAAAAATGTTTGAA 5580  
  
QY 870 AGAGGTTTACCTCTCATTCTCTAGAAAAGAGCTTAACTTCCATATCCAA 929  
DB 5581 AGAGGTTTACCTCTCATTCTCTAGAAAAGAGCTTAACTTCCATATCCAA 5640  
  
QY 930 TATTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 989  
DB 5641 TACTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700  
  
QY 990 ATATGGAATTTATTAAGAACATTTCTGCTATTTGATATTT-AGTATAAGGCAAAATAAT 1048  
DB 5701 ATATGGAATTTATTAAGAACATTTCTGCTATTTGATATTTGATATTAAGGCAAAATAAT 5760  
  
QY 1049 ATTTATGACATAACTATGAAACAGATATCTTAGGCTTTAATAACACATGATATCA 1108  
DB 5761 ATTTATGATAATAACTATGAAACAGATATCTTAGGCTTTAATAACACATGATATCA 5820  
  
QY 1109 TAAA 1112  
DB 5821 TAAA 5824

RESULT 13  
US-09-419-568F-29  
; Sequence 29, Application US/09419568F  
; Patent No. 631613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIPS) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/419,568F

; CURRENT FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; CURRENT FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 29  
; LENGTH: 5935  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-419-568F-29

Query Match 49.6%; Score 555.2; DB 3; Length 5935;  
Best Local Similarity 96.0%; Pred. No. 1.1e-129;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
  
QY 510 AAAGCTTGGAGAGAGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 569  
DB 5221 ATAGCTTGGAGAGAGCGGAGAGATCAAGCGATCGGGGAACCTGGACCTGCTGTTTATGTC 5280  
  
QY 570 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAAACGAAAGAACTGCTCCTTCCT 629  
DB 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAAACGAAAGAACTGCTCCTTCCT 5340  
  
QY 630 GCCTTCTAAAAGAACAAATAGATCCCTGATGCACTTTTCTAAGGAAAAGTGAGAA 689  
DB 5341 GCCTTCTAAAAGAACAAATAGATCCCTGATGCACTTTTCTAAGGAAAAGTGAGAA 5400  
  
QY 690 GCTAACCTGCCATCATCATTTAGAGATTTTCATGAAACCTGGCTCAGTTGAAAAGAGAAA 749  
DB 5401 GCTAACCTGCCATCATTTAGAGATTTTCATGAAACCTGGCTCAGTTGAAAAGAGAAA 5460  
  
QY 750 TAGTGTCACAGTTGTCATGAGACCCAGAGGTAGACTTGATTAACCAACAAAGATTTCATTGACA 809  
DB 5461 TAGTGTCACAGTTGTCATGAGACCCAGAGGTAGACTTGATTAACCAACAAAGATTTCATTGACA 5520  
  
QY 810 ATATTTTATGTCAGTATGATACACAGAAAATATATGTAATCTTAAAAAATGTTTGAA 869  
DB 5521 ATATTTTATGTCAGTATGATACACAGAAAATATATGTAATCTTAAAAAATGTTTGAA 5580  
  
QY 870 AGAGGTTTACCTCTCATTCTCTAGAAAAGAGCTTAACTTCCATATCCAA 929  
DB 5581 AGAGGTTTACCTCTCATTCTCTAGAAAAGAGCTTAACTTCCATATCCAA 5640  
  
QY 930 TATTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 989  
DB 5641 TACTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700  
  
QY 990 ATATGGAATTTATTAAGAACATTTCTGCTATTTGATATTT-AGTATAAGGCAAAATAAT 1048  
DB 5701 ATATGGAATTTATTAAGAACATTTCTGCTATTTGATATTTGATATTAAGGCAAAATAAT 5760  
  
QY 1049 ATTTATGACATAACTATGAAACAGATATCTTAGGCTTTAATAACACATGATATCA 1108  
DB 5761 ATTTATGATAATAACTATGAAACAGATATCTTAGGCTTTAATAACACATGATATCA 5820  
  
QY 1109 TAAA 1112  
DB 5821 TAAA 5824

RESULT 14  
US-09-354-243B-29  
; Sequence 29, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa  
; TITLE OF INVENTION: (TIPS)  
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof

FILE REFERENCE: LUD 5543.1  
CURRENT APPLICATION NUMBER: US/09/354,243B  
CURRENT FILING DATE: 1999-07-16  
PRIOR APPLICATION NUMBER: US09/178,973  
PRIOR FILING DATE: 1998-10-26  
NUMBER OF SEQ ID NOS: 29  
SEQ ID NO 29  
LENGTH: 5935  
TYPE: DNA  
ORGANISM: Homo sapiens  
FEATURE:  
US-09-354-243B-29

Query Match 49.6%; Score 555.2; DB 3; Length 5935;  
Best Local Similarity 96.0%; Pred. No. 1.1e-129;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
Qy 510 AAGCTTGGAGAGAGTGGAGAGATCAAGGGGATTTGGGAACTGACCTGCTGTTATGTC 569  
Db ATAGCTTGGAGAGAGCGAGAGATCAAGCGATCGGGAACTGACCTGCTGTTATGTC 5280  
Qy 570 TCTGAGAAATGCTTGGCTGTGAGCGAGAGAGCTAGAAAACGAAAGTCTCTCTTCT 629  
Db TCTGAGAAATGCTTGGCTGTGAGCGAGAGAGCTAGAAAACGAAAGTCTCTCTTCT 5340  
Qy 630 GCCTTCTAAAAGAACAAATAGATCCCTGATGACATCTTTTACTAAGGAAAGTGAGAA 689  
Db GCCTTCTAAAAGAACAAATAGATCCCTGATGACATCTTTTACTAAGGAAAGTGAGAA 5400  
Qy 690 GCTAACGTCCATCATCATAGAGATTTCATGAACCTGGCTCAGTTGAAAAGAAAA 749  
Db GCTAACGTCCACCATCATAGAGATTTCATGAACCTGGCTCAGTTGAAAAGAAAA 5460  
Qy 750 TAGTGTCAAGTTGTCCATGAGACGAGAGGTAGACTTGATACCAAGAAATTCATGACA 809  
Db TAGTGTCAAGTTGTCCATGAGACGAGAGGTAGACTTGATACCAAGAAATTCATGACA 5520  
Qy 810 ATATTTTATGCTACTGATGATACACAGAAAAATATGTAATTTTAAAAATTTGTTGAA 869  
Db ATATTTTATGCTACTGATGATACACAGAAAAATATGTAATTTTAAAAATTTGTTGAA 5580  
Qy 870 AGGAGTTTACCTCTCATCTCTTAAAGAAAGCTTATGTAATTTTCCATATCCAA 929  
Db AGGAGTTTACCTCTCATCTCTTAAAGAAAGCTTATGTAATTTTCCATATCCAA 5640  
Qy 930 TATTTTATATGTAAGTTTATTTATATAGTATACATTTTATTTATGTCAGTTTATTA 989  
Db TATTTTATATGTAAGTTTATTTATATAGTATACATTTTATTTATGTCAGTTTATTA 5700  
Qy 990 ATATGGATTTATTTATAGAAACATTTCTGCTATTTATTTT-AGTATAAGGCAATTAAT 1048  
Db ATATGGATTTATTTATAGAAACATTTCTGCTATTTATTTTATTTGATATAAAGCAATTAAT 5760  
Qy 1049 ATTTATGACAATACTATGGAACCAAGATATCTTAGGCTTTTAAATAACACATGATATCA 1108  
Db ATTTATGATAATACTATGGAACCAAGATATCTTAGGCTTTTAAATAACACATGATATCA 5820  
Qy 1109 TAAA 1112  
Db TAAA 5824

RESULT 15  
US-10-084-298-1  
Sequence 1, Application US/10084298  
Patent No. 6939545  
GENERAL INFORMATION:  
APPLICANT: Jacobs, Kenneth  
APPLICANT: Pittman, Debra  
APPLICANT: Fouser, Lynette  
APPLICANT: Spaulding, Vikki  
APPLICANT: Xuan, Dejun  
TITLE OF INVENTION: Composition and Method for Treating Inflammatory

TITLE OF INVENTION: Disorders  
FILE REFERENCE: G15358 CIP  
CURRENT APPLICATION NUMBER: US/10/084,298  
CURRENT FILING DATE: 2002-09-10  
PRIOR APPLICATION NUMBER: 60/270,823  
PRIOR FILING DATE: 2001-02-23  
PRIOR APPLICATION NUMBER: 60/281,353  
PRIOR FILING DATE: 2001-04-03  
PRIOR APPLICATION NUMBER: 60/131,473  
PRIOR FILING DATE: 1999-04-28  
PRIOR APPLICATION NUMBER: 09/561,811  
PRIOR FILING DATE: 2000-04-28  
NUMBER OF SEQ ID NOS: 10  
SOFTWARE: PatentIn Ver. 2.1  
SEQ ID NO 1  
LENGTH: 1191  
TYPE: DNA  
ORGANISM: Homo sapiens  
US-10-084-298-1

Query Match 47.8%; Score 535.2; DB 3; Length 1191;  
Best Local Similarity 73.4%; Pred. No. 5.7e-125;  
Matches 843; Conservative 0; Mismatches 268; Indels 37; Gaps 11;  
Qy 9 CTCTCTCTCACTTATCAACTGTGTGACATCTGTGCGATCTCTGATGGCTGTCTCTGCAGAA 68  
Db CTCTCTCTCACTTATCAACTGTGTGACATCTGTGCGATCTCTGCAATGCGCCCTGCAGAA 87  
Qy 69 ATCTATGAGTTTTCCTTATGGGACTTTGGCGCGCAGCTGCTGCTCTCTCTATGCGCCT 128  
Db ATCTATGAGTTTTCCTTATGGGACTTTGGCGCGCAGCTGCTGCTCTCTCTCTCTCTCTCT 147  
Qy 129 GTGGGCCCCAGGAGCAATGGCTGCGCCGTCACACCCGGTCAACCGGTCGAAGCTTGAGGTGTCAA 188  
Db GTGGGCCCCAGGAGCAATGGCTGCGCCGTCACACCGGTCGAAGCTTGCAAGTGTCAA 207  
Qy 189 CTTCACGACGCTGATGATGAGCAGGTCTCACTTATGCTGGCCCAAGAGGCGACCTTGC 248  
Db CTTCACGACGCTGATGATGAGCAGGTCTCACTTATGCTGGCCCAAGAGGCGACCTTGC 267  
Qy 249 AGATAACAACAAGACGCTCGGCTCATCGGGGAGAAATGTGTCGAGAGTCACTGTCTAA 308  
Db AGATAACAACAAGACGCTCGGCTCATCGGGGAGAAATGTGTCGAGAGTCACTGTCTAA 327  
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GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

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Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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3	1107.8	99.0	1166	5	US-10-084-298-3
4	1107.8	99.0	1166	6	US-10-256-977-3
5	1107.8	99.0	1166	8	US-10-873-972-3
6	1107.8	99.0	1166	10	US-11-157-387-3
7	1047.8	93.6	1111	3	US-09-751-797-9
8	1047.8	93.6	1111	7	US-10-627-273-9
9	993.2	88.8	1050	5	US-10-090-365-40
10	993.2	88.8	1050	5	US-10-104-919-42
11	993.2	88.8	1050	8	US-10-807-837-10
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13	993.2	88.8	1050	10	US-11-045-944-40
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17	601.4	53.7	7445	3	US-09-751-797-8
18	601.4	53.7	7445	7	US-10-627-273-8
19	555.2	49.6	5935	3	US-09-751-797-29
20	555.2	49.6	5935	7	US-10-627-273-29
21	535.2	47.8	1177	10	US-11-013-741-1
22	535.2	47.8	1177	10	US-11-013-920-1
23	535.2	47.8	1191	5	US-10-084-298-1

24	535.2	47.8	1191	6	US-10-256-977-1	Sequence 1, Appli
25	535.2	47.8	1191	8	US-10-873-972-1	Sequence 1, Appli
26	535.2	47.8	1191	10	US-11-157-387-1	Sequence 1, Appli
27	524.8	46.9	1152	3	US-09-870-574-1	Sequence 1, Appli
28	524.8	46.9	1152	5	US-10-006-867-153	Sequence 153, App
29	524.8	46.9	1152	5	US-10-066-500-125	Sequence 125, App
30	524.8	46.9	1152	5	US-10-063-547-153	Sequence 153, App
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37	524.8	46.9	1152	5	US-10-063-502-153	Sequence 153, App
38	524.8	46.9	1152	5	US-10-063-549-153	Sequence 243, App
39	524.8	46.9	1152	5	US-10-227-884-243	Sequence 243, App
40	524.8	46.9	1152	5	US-10-002-796-125	Sequence 125, App
41	524.8	46.9	1152	5	US-10-066-273-125	Sequence 125, App
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43	524.8	46.9	1152	5	US-10-230-163-243	Sequence 243, App
44	524.8	46.9	1152	5	US-10-063-554-153	Sequence 153, App
45	524.8	46.9	1152	5	US-10-066-269-125	Sequence 125, App

ALIGNMENTS

RESULT 1  
US-09-751-797-7  
; Sequence 7, Application US/09751797  
; Patent No. US20010024652A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/751.797  
; CURRENT FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-751-797-7

Query Match	100.0%;	Score 1119;	DB 3;	Length 1119;
Best Local Similarity	100.0%;	Pred. No. 1.1e-223;		
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Qy	121	ATTGCCCTGTGGGCCCAGGAGGCAATGCGCTGCTCAACACCCGGTGAACCTTGAG	180	
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Qy	181	GTGTCCAACTTCCAGACGCGGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGGAGGCC	240	
Db	181	GTGTCCAACTTCCAGACGCGGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGGAGGCC	240	
Qy	241	AGCCTTGCAGATAACACACAGACGTCGGCTCATCGGGAGAAACTGTTTCCGAGGAGTC	300	

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Qy 721 ATGAAACCTGCTCAGTTGAAAGAAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGA 780  
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Qy 781 GACTTGTATTAACCAACAGAGATTCATTGACAAATATTTATTTGTCATGATGATACACAGAA 840  
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Db 841 AAATAATGTACTTTTAAAGAAATGTTTGAAGAGAGTTACCTCTCATCTCTTTAGAAAAA 900  
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; Sequence 7, Application US/10627273  
; Publication No. US20040110189A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/10/627,273  
; CURRENT FILING DATE: 2003-07-25

; PRIOR APPLICATION NUMBER: US/09/751,797  
; PRIOR FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-10-627-273-7

Query Match 100.0%; Score 1119; DB 7; Length 1119;  
Best Local Similarity 100.0%; Pred. No. 1.1e-223;  
Matches 1119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 1 TAAACAGGCTCTCTCTCACTTATCAACTGTGTGACACTTGTGGATCTCTGATGGCTGTC 60  
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Qy 61 CTGCAGAAATCTATGAGTTCCTTTATGGGAGCTTTGGCGCCAGCTGCTGCTTCTC 120  
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Qy 181 GTGTCCAACTTCCAGCAGCGGTACATCGTCAACCGCACCTTTTATGTCGGCCAAAGAGGCC 240  
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RESULT 3  
US-10-084-298-3  
; Sequence 3, Application US/10084298  
; Publication No. US2003099649A1  
; GENERAL INFORMATION:  
; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: Xuan, Dejun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory  
; FILE REFERENCE: G15358 CIP  
; CURRENT APPLICATION NUMBER: US/10/084,298  
; CURRENT FILING DATE: 2002-09-10  
; PRIOR FILING DATE: 2002-09-10  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR FILING DATE: 2000-04-28  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 3  
; LENGTH: 1166  
; TYPE: DNA  
; ORGANISM: Murine

Query Match 99.0%; Score 1107.8; DB 5; Length 1166;  
Best Local Similarity 99.4%; Pred. No. 2.5e-221;  
Matches 1112; Conservative 0; Mismatches 7; Indels 0; Gaps 0;  
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Qy 121 ATTGCCCTGTGGGCCAGGAGGCAAAATGCGCTGCCGCTCAACACCCGGTCAAGCTTGAG 180  
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Db 444 AAATCAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGACCGACGAAACATCCAGAAG 503
Qy 481 AATGTCTGAGGCTGAGGAGCAGTGAAGAAAGCTTGGAGAGTGGAGAGATCAAGGCG 540
Db 504 AATGTCTGAGGCTGAGGAGCAGTGAAGAAAGCTTGGAGAGTGGAGAGATCAAGGCG 563
Qy 541 ATTGGGGAACCTGACCTGCTGTTTATGTCTCTGAGAAATGCTTGCCTCTGAGGAGAGA 600
Db 564 ATTGGGGAACCTGACCTGCTGTTTATGTCTCTGAGAAATGCTTGCCTCTGAGGAGAGA 623
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Db 684 TGGACTTTTTTACTAAAGGAAAGTGAAGCTCAAGTCCATCATCATTAGAAGATTTTAC 743
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RESULT 4  
US-10-256-977-3  
; Sequence 3, Application US/10256977  
; Publication No. US20030157106A1  
; GENERAL INFORMATION:  
; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: Xuan, Dejun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory  
; FILE REFERENCE: Disorders

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; FILE REFERENCE: GI5358 CIP
; CURRENT APPLICATION NUMBER: US/10/256,977
; CURRENT FILING DATE: 2002-09-27
; PRIOR APPLICATION NUMBER: US/10/084,298
; PRIOR FILING DATE: 2002-09-10
; PRIOR APPLICATION NUMBER: 60/270,823
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/281,353
; PRIOR FILING DATE: 2001-04-03
; PRIOR APPLICATION NUMBER: 60/131,473
; PRIOR FILING DATE: 1999-04-28
; PRIOR APPLICATION NUMBER: 09/561,811
; PRIOR FILING DATE: 2000-04-28
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn ver. 2.1
; SEQ ID NO 3
; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Murine
US-10-256-977-3

Query Match      99.0%; Score 1107.8; DB 6; Length 1166;
Best Local Similarity 99.4%; Pred. No. 2.5e-221;
Matches 1112; Conservative 0; Mismatches 7; Indels 0; Gaps 0;

Qy 1 TAAACAGGCTCTCTCTCACTTATCACTGTTGACACTTGTGCGATCTCTGATGGCTGTC 60
Db 24 TAAACAGGCTCTCTCTCAGTTATCAACTGTGTGACACTTGTGCGATCTCTGATGGCTGTC 83

Qy 61 CTGCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTGGCGCGCAGCTGCTCTCTC 120
Db 84 CTGCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTGGCGCGCAGCTGCTCTCTC 143

Qy 121 ATTGCGCTGTGGGCCAGAGGCAAAATGCGTGCCTGCTCAACACCCGGTGCAAGCTTGAG 180
Db 144 ATTGCGCTGTGGGCCAGAGGCAAAATGCGTGCCTGCTCAACACCCGGTGCAAGCTTGAG 203

Qy 181 GTGTCCAACTTCCAGCAGCGGTACATGTCGAACCGCACTTTATGCTGGCCAAAGAGGCC 240
Db 204 GTGTCCAACTTCCAGCAGCGGTACATGTCGAACCGCACTTTATGCTGGCCAAAGAGGCC 263

Qy 241 AGCCTTGAGATAAACAACAGAGCTCCGGCTCATCGGGAGAAACTGTTCCGAGGAGTC 300
Db 264 AGCCTTGAGATAAACAACAGAGTCCGGCTCATCGGGAGAAACTGTTCCGAGGAGTC 323

Qy 301 AGTGCTAAGATCAGTGCTACTGATGAAGCAGGTGCTCAACTTCACTTCCGAGAGAGCTT 360
Db 324 AGTGCTAAGATCAGTGCTACTGATGAAGCAGGTGCTCAACTTCACTTCCGAGAGAGCTT 383

Qy 361 CTGCTCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGGAGGTGGTACCTTTCTGACC 420
Db 384 CTGCTCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGGAGGTGGTACCTTTCTGACC 443

Qy 421 AAACCTCAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGACGACCAAGCAATCCAGAG 480
Db 444 AAACCTCAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGACGACCAAGCAATCCAGAG 503

Qy 481 AATGTGAGAAGCTGAAGGAGACAGTGAAGAAAGCTTGGAGAGAGTGGAGAGATCAAGGCG 540
Db 504 AATGTGAGAAGCTGAAGGAGACAGTGAAGAAAGCTTGGAGAGAGTGGAGAGATCAAGGCG 563

Qy 541 ATTGGGGAACCTGGACCTCTGTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGA 600
Db 564 ATTGGGGAACCTGGACCTCTGTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGA 623

Qy 601 AGCTAGAAAAACGAAGAACTGCTCTCTGCTCTCTTAAAGAAACAAATAAGATCCCTGAA 660
Db 624 AGCTAGAAAAACGAAGAACTGCTCTCTCTGCTCTCTTAAAGAAACAAATAAGATCCCTGAA 683

Qy 661 TGGACTTTTTTACTAAAGAAAGTGAAGAGCTTAACGTCATCATCATTTAGAAAGATTTCAC 720
Db 684 TGGACTTTTTTACTAAAGAAAGTGAAGAGCTTAACGTCATCATCATTTAGAAAGATTTCAC 743

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Qy 721 ATGAAACCTGGCTCAGTTGAAAAAGAAAAATAGTGTCAAGTTGTCCATGAGACCAAGGTA 780
Db 744 ATGAAACCTGGCTCAGTTGAAAAAGAAAAATAGTGTCAAGTTGTCCATGAGACCAAGGTA 803

Qy 781 GACTTGATTAACCAAAAGATTCTTGTGACATATTTTATGCTACTGATGATCAACAGAA 840
Db 804 GACTTGATTAACCAAAAGATTCTTGTGACATATTTTATGCTACTGATGATCAACAGAA 863

Qy 841 AATATATGTACTTTTAAAAAATGTTTGAAGAGAGGTGTACCTCTCATTCCTTTAGAAAAA 900
Db 864 AATATATGTACTTTTAAAAAATGTTTGAAGAGAGGTGTACCTCTCATTCCTTTAGAAAAA 923

Qy 901 AGCTTATGTAACTTCAATTTCCATATCCAATATTTTATATATGTAAGTTTATTTATATA 960
Db 924 AGCTTATGTAACTTCAATTTCCATATCCAATATTTTATATATGTAAGTTTATTTATTATA 983

Qy 961 GTATACATTTTATATGTCAGTTTATATATGATGATTTTATATAGAAAAATTTATCTGC 1020
Db 984 GTATACATTTTATATGTCAGTTTATATATGATGATTTTATATAGAAAAATTTATCTGC 1043

Qy 1021 TATTGATATTTAGTATAAGGCAATATATTTTATGACAAATACTATGAAAAACAAGATATC 1080
Db 1044 TATTGATATTTAGTATAAGGCAATATATTTTATGACAAATACTATGAAAAACAAGATATC 1103

Qy 1081 TTAGGCTTTAATAAACAACATGGATATCATAAAAA 1119
Db 1104 TTAGGCTTTAATAAACAACATGGATATCATAAAAA 1142

RESULT 5
US-10-873-972-3
; Sequence 3, Application US/10873972
; Publication No. US2005004220A1
; GENERAL INFORMATION:
; APPLICANT: Li, Jing
; APPLICANT: Tan, Xiang-Yang
; APPLICANT: Tomkinson, Kathleen N.
; APPLICANT: Pittman, Debra D.
; APPLICANT: Veldman, Geertruida M.
; APPLICANT: Fouser, Lynette
; TITLE OF INVENTION: Antibodies Against Interleukin-22 and Uses Therefor
; FILE REFERENCE: AM101524
; CURRENT APPLICATION NUMBER: US/10/873,972
; CURRENT FILING DATE: 2004-06-22
; PRIOR APPLICATION NUMBER: US 60/480,652
; PRIOR FILING DATE: 2003-06-23
; PRIOR APPLICATION NUMBER: US 10/084,298
; PRIOR FILING DATE: 2002-02-25
; PRIOR APPLICATION NUMBER: US 60/270,823
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: US 60/281,353
; PRIOR FILING DATE: 2001-04-03
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Mus musculus
US-10-873-972-3

Query Match      99.0%; Score 1107.8; DB 8; Length 1166;
Best Local Similarity 99.4%; Pred. No. 2.5e-221;
Matches 1112; Conservative 0; Mismatches 7; Indels 0; Gaps 0;

Qy 1 TAAACAGGCTCTCTCTCACTTATCAACTGTTGACACTTGTGCGATCTCTGATGGCTGTC 60
Db 24 TAAACAGGCTCTCTCTCAGTTATCAACTGTGTGACACTTGTGCGATCTCTGATGGCTGTC 83

Qy 61 CTGCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTGGCGCGCAGCTGCTCTCTC 120
Db 84 CTGCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTGGCGCGCAGCTGCTCTCTC 143

Qy 121 ATTGCGCTGTGGGCCAGAGGCAAAATGCGTGCCTGCTCAACACCCGGTGCAAGCTTGAG 180

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Db 144 ATTGCCCCGTGGGCCCCAGGAGGCAAAATGCGCTGCCCGTCAACACCCGGTCAAGAGCTTGAG 203
Qy 181 GTGTCCAACTTCCAGACGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGAGGCC 240
Db 204 GTGTCCAACTTCCAGACGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGAGGCC 263
Qy 241 AGCTTTCAGATACAAACAGACGCTCCGCTCATCGGGGAGAAACTGTTCCGAGAGTC 300
Db 264 AGCTTTCAGATACAAACAGACGCTCCGCTCATCGGGGAGAAACTGTTCCGAGAGTC 323
Qy 301 AGTGCTAAAGATCAGTGCTACCTGATGAACAGAGTGTCAACTCACCCTGGAGAGCTT 360
Db 324 AGTGCTAAAGATCAGTGCTACCTGATGAACAGAGTGTCAACTCACCCTGGAGAGCTT 383
Qy 361 CTGCTCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGAGGTGGTACCTTTCCTGACC 420
Db 384 CTGCTCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGAGGTGGTACCTTTCCTGACC 443
Qy 421 AAATCAGCAATCAGCTCAGCTCCTGTGTCATCAGCGGTGACGACCAAGAAATCCAGAG 480
Db 444 AAATCAGCAATCAGCTCAGCTCCTGTGTCATCAGCGGTGACGACCAAGAAATCCAGAG 503
Qy 481 AATGTCAGAGGCTGAGGAGACAGTGAAGAAAGCTTGGAGAGTGGAGAGATCAAGGCG 540
Db 504 AATGTCAGAGGCTGAGGAGACAGTGAAGAAAGCTTGGAGAGTGGAGAGATCAAGGCG 563
Qy 541 ATTGGGGAATCGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCGTCTGAGCGAGAGA 600
Db 564 ATTGGGGAATCGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCGTCTGAGCGAGAGA 623
Qy 601 AGCTAGAAAACGAAAGACTGCTCCTCCCTCTTAAAGAAACAAATAGATCCCTGAA 660
Db 624 AGCTAGAAAACGAAAGACTGCTCCTCCCTCTTAAAGAAACAAATAGATCCCTGAA 683
Qy 661 TGGACTTTTTTACTAAAGGAAAGTGAAGCTCAAGCTCAATCATCATTAGAAGATTTCAC 720
Db 684 TGGACTTTTTTACTAAAGGAAAGTGAAGCTCAAGCTCAATCATCATTAGAAGATTTCAC 743
Qy 721 ATGAAACCTGGCTCAGTTGAAAGAAATAGTGTCAAGTTGTCATGAGACCAAGAGTA 780
Db 744 ATGAAACCTGGCTCAGTTGAAAGAAATAGTGTCAAGTTGTCATGAGACCAAGAGTA 803
Qy 781 GACTTGATAACCAAGATTCATGTACAATATTTTATGCTCACTGATGATACAAAGAA 840
Db 804 GACTTGATAACCAAGATTCATGTACAATATTTTATGCTCACTGATGATACAAAGAA 863
Qy 841 AAATAATGTAATTTAAAGAAATTTGTTGAAAGAGGTTTACCTCTCATTTCTTTAGAAAAA 900
Db 864 AAATAATGTAATTTAAAGAAATTTGTTGAAAGAGGTTTACCTCTCATTTCTTTAGAAAAA 923
Qy 901 AGCTTATGTAATTCATTTCCATATCCAATATTTTATATATGTAAGTTTATTTATTATAA 960
Db 924 AGCTTATGTAATTCATTTCCATATCCAATATTTTATATATGTAAGTTTATTTATTATAA 983
Qy 961 GTATACATTTTATTTATGTCAGTTTATTAATATGAGTTTATTTATAGAAACATTATCTGC 1020
Db 984 GTATACATTTTATTTATGTCAGTTTATTAATATGAGTTTATTTATAGAAACATTATCTGC 1043
Qy 1021 TATTGATATTTATGTAAGGCAATATATTTATGACAAATATGAGAAACAAAGATATC 1080
Db 1044 TATTGATATTTATGTAAGGCAATATATTTATGACAAATATGAGAAACAAAGATATC 1103
Qy 1081 TTAGGCTTTTAAACACATGATATCATATAAAAAA 1119
Db 1104 TTAGGCTTTTAAACACATGATATCATATAAAAAA 1142
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## RESULT 6

US-11-157-387-3

; Sequence 3, Application US/11157387

; Publication No. US20050238649A1

; GENERAL INFORMATION:

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; APPLICANT: Jacobs, Kenneth
; APPLICANT: Pittman, Debra
; APPLICANT: Fouser, Lynette
; APPLICANT: Spaulding, Vikki
; APPLICANT: Xuan, Dejun
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory
; TITLE OF INVENTION: Disorders
; FILE REFERENCE: G15358 CIP
; CURRENT APPLICATION NUMBER: US/11/157,387
; CURRENT FILING DATE: 2005-06-20
; PRIOR APPLICATION NUMBER: US/10/084,298
; PRIOR FILING DATE: 2002-09-10
; PRIOR APPLICATION NUMBER: 60/270,823
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/281,353
; PRIOR FILING DATE: 2001-04-03
; PRIOR APPLICATION NUMBER: 60/131,473
; PRIOR FILING DATE: 1999-04-28
; PRIOR APPLICATION NUMBER: 09/561,811
; PRIOR FILING DATE: 2000-04-28
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Murine
; US-11-157-387-3
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Query Match 99.0%; Score 1107.8; DB 10; Length 1166;
Best Local Similarity 99.4%; Pred. No. 2.5e-221;
Matches 1112; Conservative 0; Mismatches 7; Indels 0; Gaps 0;

Qy 1 TAAACAGGCTCTCTCTCTCACTTATCAACTGTGACACTTGTGGGATCTCTGATGGCTGTC 60
Db 24 TAAACAGGCTCTCTCTCTCACTTATCAACTGTGACACTTGTGGGATCTCTGATGGCTGTC 83
Qy 61 CTGCAAGAAATCTATGATGTTTTCCTTTATGGGACATTTGGCCCGCAGCTGCTGCTTCTC 120
Db 84 CTGCAAGAAATCTATGATGTTTTCCTTTATGGGACATTTGGCCCGCAGCTGCTGCTTCTC 143
Qy 121 ATTGCCCTGTGGCCCGCAGGAGGCAAAATGGCTCGCCGTCAACACCCGGTCAAGAGCTTGAG 180
Db 144 ATTGCCCTGTGGCCCGCAGGAGGCAAAATGGCTCGCCGTCAACACCCGGTCAAGAGCTTGAG 203
Qy 181 GTGTCCAACTTCCAGACGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGAGGCC 240
Db 204 GTGTCCAACTTCCAGACGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGAGGCC 263
Qy 241 AGCTTTCAGATACAAACAGACGCTCCGCTCATCGGGGAGAAACTGTTCCGAGAGTC 300
Db 264 AGCTTTCAGATACAAACAGACGCTCCGCTCATCGGGGAGAAACTGTTCCGAGAGTC 323
Qy 301 AGTGCTAAAGATCAGTGCTACCTGATGAACAGAGTGTCAACTCACCCTGGAGAGCTT 360
Db 324 AGTGCTAAAGATCAGTGCTACCTGATGAACAGAGTGTCAACTCACCCTGGAGAGCTT 383
Qy 361 CTGCTCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGAGGTGGTACCTTTCCTGACC 420
Db 384 CTGCTCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGAGGTGGTACCTTTCCTGACC 443
Qy 421 AAATCAGCAATCAGCTCAGCTCCTGTGTCATCAGCGGTGACGACCAAGAAATCCAGAG 480
Db 444 AAATCAGCAATCAGCTCAGCTCCTGTGTCATCAGCGGTGACGACCAAGAAATCCAGAG 503
Qy 481 AATGTCAGAGGCTGAGGAGACAGTGAAGAAAGCTTGGAGAGTGGAGAGATCAAGGCG 540
Db 504 AATGTCAGAGGCTGAGGAGACAGTGAAGAAAGCTTGGAGAGTGGAGAGATCAAGGCG 563
Qy 541 ATTGGGGAATCGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCGTCTGAGCGAGAGA 600
Db 564 ATTGGGGAATCGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCGTCTGAGCGAGAGA 623
Qy 601 AGCTAGAAAACGAAAGACTGCTCCTCCCTCTTAAAGAAACAAATAGATCCCTGAA 660
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Db	624	AGCTAGAAAACGAAGAACTGCTCTTCTGCTCTTCTTAAAGAAACAATAAGATCCCTGAA	683	
Qy	661	TGGACTTTTCTAAAGGAAAGTGAGAACTAAGCTTCCATCATCATAGATTTTAC	720	
Db	684	TGGACTTTTCTAAAGGAAAGTGAGAACTAAGCTTCCATCATCATAGATTTTAC	743	
Qy	721	ATGAAACCTGGCTCAGTTGAAAAGAAAATAGTCAAGTTGTCATGAGACCAAGAGTA	780	
Db	744	ATGAAACCTGGCTCAGTTGAAAAGAAAATAGTCAAGTTGTCATGAGACCAAGAGTA	803	
Qy	781	GACTTTGATAACCAACAAAGATTCAATGACAAATATTTTATGTCATGATGATACACAGAA	840	
Db	804	GACTTTGATAACCAACAAAGATTCAATGACAAATATTTTATGTCATGATGATACACAGAA	863	
Qy	841	AAATAATGTAACCTTTAAAGGAAAGTTTGAAGAGGAGTTACCTCTCATTTGAGAAAAA	900	
Db	864	AAATAATGTAACCTTTAAAGGAAAGTTTGAAGAGGAGTTACCTCTCATTTGAGAAAAA	923	
Qy	901	AGCTTATGTAACCTTTTCCATATCCAAATATTTTATATGTAAGTTTATTTATTATATA	960	
Db	924	AGCTTATGTAACCTTTTCCATATCCAAATATTTTATATGTAAGTTTATTTATTATATA	983	
Qy	961	GTATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAACATTTATCTGC	1020	
Db	984	GTATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAACATTTATCTGC	1043	
Qy	1021	TATTGATATTTAGTATAGGCAATATATTTATGACAAATACATGGAACCAAGATATC	1080	
Db	1044	TATTGATATTTAGTATAGGCAATATATTTATGACAAATACATGGAACCAAGATATC	1103	
Qy	1081	TTAGGCTTTTATATAACACATGATATCATATAAAAAA	1119	
Db	1104	TTAGGCTTTTATATAACACATGATATCATATAAAAAA	1142	
RESULT 7				
US-09-751-797-9				
; Sequence 9, Application US/09751797				
; Patent No. US20010024652A1				
; GENERAL INFORMATION:				
; APPLICANT: Dumoutier, Laure				
; APPLICANT: Renauld, Jean-Christophe				
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac				
; TITLE OF INVENTION: (TIFs) The Proteins Encoded, and Uses Thereof				
; FILE REFERENCE: LUD 5543.2				
; CURRENT APPLICATION NUMBER: US/09/751,797				
; CURRENT FILING DATE: 2000-12-29				
; PRIOR APPLICATION NUMBER: 09/419,568				
; PRIOR FILING DATE: 1999-10-18				
; PRIOR APPLICATION NUMBER: US09/178,973				
; PRIOR FILING DATE: 1998-10-26				
; NUMBER OF SEQ ID NOS: 29				
; SEQ ID NO 9				
; LENGTH: 1111				
; TYPE: DNA				
; ORGANISM: Mus musculus				
; FEATURE:				
US-09-751-797-9				
Query Match 93.6%; Score 1047.8; DB 3; Length 1111;				
Best Local Similarity 97.0%; Pred. No. 8.6e-209;				
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;				
Qy	3	AACGGCTCTCTCTCACTTATCACTTGTGACACTTGTGGGATCTCTGTGCTGCTGCT	62	
Db	1	AACGGCTCTCTCTCACTTATCACTTGTGACACTTGTGGGATCTCTGTGCTGCTGCT	60	
Qy	63	GCAGAAATCTATGAGTTTTTCCCTTATGGGACATTTGGCCGCGAGCTGCTTCTCAT	122	
Db	61	GCAGAAATCTATGAGTTTTTCCCTTATGGGACATTTGGCCGCGAGCTGCTTCTCAT	120	

RESULT 8  
US-10-627-273-9  
; Sequence 9, Application US/10627273  
; Publication No. US20040110189A1

Qy	123	TGCCCTGTGGGCCCCAGAGGCAAAATGCGTCCCGTCAACACCCGGTGCAGCTTGAAGT	182	
Db	121	TGCCCTGTGGGCCCCAGAGGCAAAATGCGTCCCGTCAACACCCGGTGCAGCTTGAAGT	180	
Qy	183	GTCAAACTTCCAGAGCCGTACATGTCACACCCGACCTTTATGCTGCGCAAGGAGCCAG	242	
Db	181	GTCAAACTTCCAGAGCCGTACATGTCACACCCGACCTTTATGCTGCGCAAGGAGCCAG	240	
Qy	243	CCTTGCAGATAACAAACACAGACGTCGGGCTCATCGGGGAGAAACTGTTCCGAGAGTCAG	302	
Db	241	CCTTGCAGATAACAAACACAGACGTCGGGCTCATCGGGGAGAAACTGTTCCGAGAGTCAG	300	
Qy	303	TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCCCTCGAAGACGTTCT	362	
Db	301	TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCCCTCGAAGACATTTCT	360	
Qy	363	GCTCCCCCAGTCAGACAGGTTCCAGCCCTACATGTCAGGAGGTGGTACTTTTCTTGACAA	422	
Db	361	GCTCCCCCAGTCAGACAGGTTCCAGCCCTACATGTCAGGAGGTGGTGGCTTTTCTTGACAA	420	
Qy	423	ACTCAGCAATCAGCTCAGCTCCTGTGCATCAGCGGTGACGACCAGAAACATCCAGAAAGAA	482	
Db	421	ACTCAGCAATCAGCTCAGCTCCTGTGCATCAGTGTGACGACCAAGAAACATCCAGAAAGAA	480	
Qy	483	TGTCAGAAAGCTGAAGAGGACAGTGAAGAAAGCTTTGGAGAGAGTGGAGAGATCAAGGCGAT	542	
Db	481	TGTCAGAAAGCTGAAGAGGACAGTGAAGAAAGCTTTGGAGAGAGTGGAGAGATCAAGGCGAT	540	
Qy	543	TGGGAACTCGACCTGCTGTTTATGCTCTGAGAAATGCTTGGTCTGAGGAGAGAGAG	602	
Db	541	CGGGAACTCGACCTGCTGTTTATGCTCTGAGAAATGCTTGGTCTGAGGAGAGAGAG	600	
Qy	603	CTAGAAAACGAAGAACTGCTCTTCTGCGCTTCTTAAAGAAACAAATGAATCCCTGGAATG	662	
Db	601	CTAGAAAACGAAGAACTGCTCTTCTGCGCTTCTTAAAGAAACAAATGAATCCCTGGAATG	660	
Qy	663	GACTTTTTTACTTAAAGGAAAGTGAGAAAGCTAAACGTCATCATCATTTAGAAAGATTTCAAT	722	
Db	661	GACTTTTTTACTTAAAGGAAAGTGAGAAAGCTAAACGTCATCATCATTTAGAAAGATTTCAAT	720	
Qy	723	GAAACCTGGCTCAGTTGAAAAGAAAGAAATAGTGTCAAGTTGTCATGACAGAGGTAGA	782	
Db	721	GAAACCTGGCTCAGTTGAAAAGAGAAAAATAGTGTCAAGTTGTCATGACAGAGGTAGA	780	
Qy	783	CTTGATAACCAAGAAATTCATTTGACAAATATTTTATTTGTCATGATGATCAACAGAAAA	842	
Db	781	CTTGATAACCAAGAAATTCATTTGACAAATATTTTATTTGTCATGATGATGCAACAGAAAA	840	
Qy	843	ATAATGTACTTTTAAAAAATTTGTTTGAAGAGGAGTTACCTCTCAATTCCTTTAGAAAAAAG	902	
Db	841	AGTATGTACTTTTAAAAAATTTGTTTGAAGAGGAGTTACCTCTCAATTCCTTAGAAAAAAG	900	
Qy	903	CTTATGTAACTTCAATTTCCATATCCAAATATTTTATATATGTAAGTTTATTTATTAAGT	962	
Db	901	CCTATGTAACTTCAATTTCCATATCCAAATATTTTATATATGTAAGTTTATTTATTAAGT	960	
Qy	963	ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAACATTTATCTGCTA	1022	
Db	961	ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAACATTTATCTGATG	1020	
Qy	1023	TTGATATTTT-AGTATAAGGCAAAATAATATTTTATGACAAATACATATGGAACCAAGATATCT	1081	
Db	1021	TTGATATTTTGGTATATTAAGCAAAATAATATTTATGATAATACTATAGAAACCAAGATATCT	1080	
Qy	1082	TAGGCTTTTAAATAACACATGATATCATATA	1112	
Db	1081	TAGGCTTTTAAATAACACATGATATCATATA	1111	

GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louheid, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Protein  
; TITLE OF INVENTION: (TIPs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/10/627,273  
; PRIOR FILING DATE: 2003-07-25  
; CURRENT APPLICATION NUMBER: US/09/751,797  
; PRIOR FILING DATE: 2000-12-29  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-10-627-273-9

Query Match  
Best Local Similarity 97.0%; Pred. No. 8.6e-209;  
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;

3 AACAGGCTCTCCCTCACTTATCAACTTGTGACACTTGTGCGATCTCTGATGGCTGTCCT 62  
1 AACAGGCTCTCCCTCACTTATCAACTTGTGACACTTGTGCGATCTCTGATGGCTGTCCT 60

63 GCAGAAATCTATGAGTTTTTCCCTATGAGGACTTTTGGCCGCGAGCTGCTGCTTCTCAT 122  
61 GCAGAAATCTATGAGTTTTTCCCTATGAGGACTTTTGGCCGCGAGCTGCTGCTTCTCAT 120

123 TGGCTGTGGCCGAGAGGCAATGCGCTGCGGTCAACACCGGCTGCAAGCTTGAGGT 182  
121 TGGCTGTGGCCGAGAGGCAATGCGCTGCGGTCAACACCGGCTGCAAGCTTGAGGT 180

183 GTCCAACTTCCAGAGCGGTACATCGTCAACCGGCTTATGCTGCGCAAGAGGCGAG 242  
181 GTCCAACTTCCAGAGCGGTACATCGTCAACCGGCTTATGCTGCGCAAGAGGCGAG 240

243 CCTTGACAGATAACACACACAGAGCTCGGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 302  
241 CCTTGACAGATAACACACACAGAGCTCGGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 300

303 TGCTAAAGATCAGTGTCTACCTGATGAAGAGAGGTGCTCAACTTCACTTGGAAAGAGTCT 362  
301 TGCTAAAGATCAGTGTCTACCTGATGAAGAGAGGTGCTCAACTTCACTTGGAAAGAGTCT 360

363 GCTCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGAGGTGCTTCTTCTGACCA 422  
361 GCTCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGAGGTGCTTCTTCTGACCA 420

423 ACTCAGCAATCAGCTCAGCTCCTGTGTCATCATGCGGTGACGACACAGAACTCCAGAGAA 482  
421 ACTCAGCAATCAGCTCAGCTCCTGTGTCATCATGCGGTGACGACACAGAACTCCAGAGAA 480

483 TGTGAGAGGCTGAAGAGAGCAGTGAAGAAAGCTTGGAGAGAGTGGAGAGATCAAGGCGAT 542  
481 TGTGAGAGGCTGAAGAGAGCAGTGAAGAAAGCTTGGAGAGAGTGGAGAGATCAAGGCGAT 540

543 TGGGGAATGGAAGCTGCTGTTATGCTGTCAGAAATGCTTGGCTGTCGAGGAGAGAG 602  
541 CGGGGAATGGAAGCTGCTGTTATGCTGTCAGAAATGCTTGGCTGTCGAGGAGAGAG 600

603 CTAGAAACGAGAACTGCTTCTTCTGCTTCTTAAAGAAACAAATGAAGATCCCTGAATG 662  
601 CTAGAAACGAGAACTGCTTCTTCTGCTTCTTAAAGAAACAAATGAAGATCCCTGAATG 660

663 GACTTTTTTCTAAGGAAGTGAAGAGCTTAAAGCTTCAATCATTTAGAGAGATTTTCAAT 722

Db 661 GACTTTTTTCTAAGGAAGTGAAGAGCTTAAAGCTTCAATCATTTAGAAAGATTTTCAAT 720  
Qy 723 GAAACCTGGCTCAGTTTCAAAAAGAAATAGTGTCAAGTTGTCTCCATGAGACCCAGAGGTAGA 782  
Db 721 GAAACCTGGCTCAGTTTCAAAAAGAAATAGTGTCAAGTTGTCTCCATGAGACCCAGAGGTAGA 780  
Qy 783 CTTGATAACCAAGAAATTCATTGACAAATATTTTATTTGTCACATGATGATCAACAGAAAA 842  
Db 781 CTTGATAACCAAGAAATTCATTGACAAATATTTTATTTGTCATTAATGATCAACAGAAAA 840  
Qy 843 ATAATGCTACTTTTAAAAAATTTGTTGAAAGGAGGTACCTCTCATTTCTCTTAGAAAAAAG 902  
Db 841 AGTATGCTACTTTTAAAAAATTTGTTGAAAGGAGGTACCTCTCATTTCTCTTAGAAAAAAG 900  
Qy 903 CTTATGTAACCTTCAATTTCCATATFCCAATATTTTATATATGTAAGTTTATTTATTAAGT 962  
Db 901 CCTATGTAACCTTCAATTTCCATATFCCAATATTTTATATATGTAAGTTTATTTATTAAGT 960  
Qy 963 ATACATTTTATTTATGCTAGTTTATTAATATGAGTTTATTTATAGAAAAATTTATCTCTA 1022  
Db 961 ATACATTTTATTTATGCTAGTTTATTAATATGAGTTTATTTATAGAAAAATTTATCTGATG 1020  
Qy 1023 TTGATATTT-AGTATAGGCAATAATATTTTATGCAATAACTATGGAACAAGATATCT 1081  
Db 1021 TTGATATTTGAGTATTAAGCAATAATATTTTATGTAATAACTATAGAAAAAGATATCT 1080  
Qy 1082 TAGGCTTTTAAATAACACATGGATATCATAAA 1112  
Db 1081 TAGGCTTTTAAATAACACATGAATATCATAAA 1111

## RESULT 9

US-10-090-365-40  
; Sequence 40, Application US/10090365  
; Publication No. US20030077706A1  
; GENERAL INFORMATION:  
; APPLICANT: Presnell, Scott R.  
; APPLICANT: Xu, Wenfeng  
; APPLICANT: Kindsvogel, Wayne  
; APPLICANT: Chen, Zhi  
; TITLE OF INVENTION: Mouse Cytokine Receptor  
; FILE REFERENCE: 01-08  
; CURRENT APPLICATION NUMBER: US/10/090,365  
; PRIOR FILING DATE: 2002-03-04  
; PRIOR APPLICATION NUMBER: US 60/273,035  
; PRIOR FILING DATE: 2001-03-02  
; PRIOR APPLICATION NUMBER: US 60/279,232  
; PRIOR FILING DATE: 2001-03-27  
; NUMBER OF SEQ ID NOS: 49  
; SOFTWARE: FastSeq for Windows Version 3.0  
; SEQ ID NO 40  
; LENGTH: 1050  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (50)...(589)  
US-10-090-365-40

Query Match 88.8%; Score 993.2; DB 5; Length 1050;  
Best Local Similarity 97.2%; Pred. No. 2.2e-197;  
Matches 1021; Conservative 0; Mismatches 28; Indels 1; Gaps 1;

Qy 3 AACAGGCTCTCCCTCACTTCACTTCACTTGTGACACTTGTGCGATCTCTGATGGCTGTCCT 62  
Db 1 AACAGGCTCTCCCTCACTTCACTTCACTTGTGACACTTGTGCGATCGGTGATGGCTGTCCT 60

Qy 63 GCAGAAATCTATGAGTTTTTCCCTTATGAGGAACTTTTGGCCGCGAGCTGCTGCTTCTCAT 122  
Db 61 GCAGAAATCTATGAGTTTTTCCCTTATGAGGAACTTTTGGCCGCGAGCTGCTGCTTCTCAT 120

Qy 123 TGGCTGTGGCCGAGAGGCAATGCGCTGCGGTCAACACCGGCTGCAAGCTTGAGGT 182

Db 121 TGCCCTGTGGCCCCAGGAGGCAAAATGCGCTGCCCATCAACACCCGGTGCAAGCTTGAGGT 180  
Qy 183 GTCCAACTTCAGACAGCGGTACATCGTCAACCGCACCTTTATGTCGGCCAGAGGGCCAG 242  
Db 181 GTCCAACTTCAGACAGCGGTACATCGTCAACCGCACCTTTATGTCGGCCAGAGGGCCAG 240  
Qy 243 CTTTGACAGATAACAACACAGACGTCGGCTCATCGGGAGAGAACTGTTCCGAGAGTCCAG 302  
Db 241 CTTTGACAGATAACAACACAGACGTCGGCTCATCGGGAGAGAACTGTTCCGAGAGTCCAG 300  
Qy 303 TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCAACCTTGAAGACGTTCT 362  
Db 301 TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCAACCTTGAAGACATTTCT 360  
Qy 363 GCTCCCCAGTCACACAGGTTCAGCGCTTCCAGCTCATCATCGAGAGGTGTACTTTCCTGACCA 422  
Db 361 GCTCCCCAGTCACACAGGTTCAGCGCTTCCAGCTCATCATCGAGAGGTGTACTTTCCTGACCA 420  
Qy 423 ACTCAGCAATCAGTGTCTGCTCATCATCAGCGGTGACGACCAAGAACATCCAGAGAGAA 482  
Db 421 ACTCAGCAATCAGTGTCTGCTCATCATCAGTGTGTCGACCAAGAACATCCAGAGAGAA 480  
Qy 483 TGTGAGAGGCTGAAGGAGACAGTGAAGAAAGCTTTGGAGAGAGTGGAGAGATCAAGGCGAT 542  
Db 481 TGTGAGAGGCTGAAGGAGACAGTGAAGAAAGCTTTGGAGAGAGCGGAGAGATCAAGGCGAT 540  
Qy 543 TGGGGAAGTGAACCTGCTGTTTATGTCCTCAGAGAAATGCTTGGCTCTGAGCGAGAGAG 602  
Db 541 CGGGGAAGTGAACCTGCTGTTTATGTCCTCAGAGAAATGCTTGGCTCTGAGCGAGAGAG 600  
Qy 603 CTAGAGAAAGAGAACTGCTCTCTCTGCTTCTTAAAGAAAGAACTAAGATCCCTGAAATG 662  
Db 601 CTAGAGAAAGAGAACTGCTCTCTCTGCTTCTTAAAGAAAGAACTAAGATCCCTGAAATG 660  
Qy 663 GACTTTTTTACTAAGGAAAGTGAGAGCTAACGTCATCATCATTTAGAGAGATTTCAAT 722  
Db 661 GACTTTTTTACTAAGGAAAGTGAGAGCTAACGTCATCATCATTTAGAGAGATTTCAAT 720  
Qy 723 GAAACCTGGCTCAGTTGAAAAGAAATAGTGTCAAGTGTCCATGATGACACAGAGGTAGA 782  
Db 721 GAAACCTGGCTCAGTTGAAAAGAAATAGTGTCAAGTGTCCATGATGACACAGAGGTAGA 780  
Qy 783 CTTGATACCAACAAGATTTCATGACATAATTTTATGTCACCTGATGATACACAGAGAAA 842  
Db 781 CTTGATACCAACAAGATTTCATGACATAATTTTATGTCATTTGATGATGACAGAGAAA 840  
Qy 843 ATATATGTAATTTAAATAATGTTTGAAGGAGGTACCTCTCATTCCTTTAGAGAAAAAG 902  
Db 841 AGTATGTAATTTAAATAATGTTTGAAGGAGGTACCTCTCATTCCTCTAGAGAGAAAAG 900  
Qy 903 CTTATGTAATTTCAATTCCTCATATCCATATTTTATATATATGTAAGTTTATTTATTAAGT 962  
Db 901 CCTATGTAATTTCAATTCCTCATATCCATATTTTATATATGTAAGTTTATTTATTAAGT 960  
Qy 963 ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAACATTTATCTGCTA 1022  
Db 961 ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAAAATTTATCTGATG 1020  
Qy 1023 TTGATATTT-AGTATAAGGCAATAATATT 1051  
Db 1021 TTGATATTTGATATAAGCAATAATATT 1050

RESULT 10

US-10-104-919-42  
; Sequence 42, Application US/10104919  
; Publication No. US20030099608A1  
; GENERAL INFORMATION:  
; APPLICANT: Fresnell, Scott R.  
; APPLICANT: Xu, Wenfeng  
; APPLICANT: Kindsvogel, Wayne  
; APPLICANT: Chen, Zhi  
; APPLICANT: Hughes, Steven D.

; TITLE OF INVENTION: Human Cytokine Receptor  
; FILE REFERENCE: 01-12  
; CURRENT APPLICATION NUMBER: US/10/104,919  
; CURRENT FILING DATE: 2002-03-23  
; PRIOR APPLICATION NUMBER: US 60/279,222  
; PRIOR FILING DATE: 2001-03-27  
; NUMBER OF SEQ ID NOS: 62  
; SOFTWARE: Fast-Seq for Windows Version 3.0  
; SEQ ID NO 42  
; LENGTH: 1050  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (5)...(589)  
US-10-104-919-42

Query Match 88.8%; Score 993.2; DB 5; Length 1050;  
Best Local Similarity 97.2%; Pred. No. 2.2e-197;  
Matches 1021; Conservative 0; Mismatches 28; Indels 1; Gaps 1;

Qy 3 AACAGGCTCTCCTCTCACTTATCAACTGTTGACACTTTGTGGATCTCTGATGGCTGTCT 62  
Db 1 AACAGGCTCTCCTCTCACTTATCAACTTTTGCATCTTGTGCGATCGGTGATGGCTGTCT 60  
Qy 63 GCAGAAATCTATGAGTGTTCCTTATGCGGACTTTTGGCCGCCAGCTGCTGCTTCTCAT 122  
Db 61 GCAGAAATCTATGAGTGTTCCTTATGCGGACTTTTGGCCGCCAGCTGCTGCTTCTCAT 120  
Qy 123 TGCCCTGTGGCCCCAGGAGCAAAATGCGTCCCGCTCAACACCCGGTGCAAGCTTGAGGT 182  
Db 121 TGCCCTGTGGCCCCAGGAGCAAAATGCGTCCCGCTCAACACCCGGTGCAAGCTTGAGGT 180  
Qy 183 GTCCAACTTCAGACAGCGGTACATCGTCAACCGCACCTTTATGTCGGCCAGAGGGCCAG 242  
Db 181 GTCCAACTTCAGACAGCGGTACATCGTCAACCGCACCTTTATGTCGGCCAGAGGGCCAG 240  
Qy 243 CTTTGACAGATAACAACACAGACGTCGGCTCATCGGGAGAGAACTGTTCCGAGAGGTGAG 302  
Db 241 CTTTGACAGATAACAACACAGACGTCGGCTCATCGGGAGAGAACTGTTCCGAGAGGTGAG 300  
Qy 303 TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCAACCTTGAAGACGTTCT 362  
Db 301 TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCAACCTTGAAGACATTTCT 360  
Qy 363 GCTCCCCAGTCACACAGGTTCAGCGCTCATCATCGAGAGGTGTACTTTCCTGACCAA 422  
Db 361 GCTCCCCAGTCACACAGGTTCAGCGCTCATCATCGAGAGGTGTGCTTTCCTGACCAA 420  
Qy 423 ACTCAGCAATCAGTGTCTGCTCATCATCAGCGGTGACGACCAAGAACATCCAGAGAGAA 482  
Db 421 ACTCAGCAATCAGTGTCTGCTCATCATCAGTGTGTCGACCAAGAACATCCAGAGAGAA 480  
Qy 483 TGTGAGAGGCTGAAGGAGACAGTGAAGAAAGCTTTGGAGAGAGTGGAGAGATCAAGGCGAT 542  
Db 481 TGTGAGAGGCTGAAGGAGACAGTGAAGAAAGCTTTGGAGAGAGCGGAGAGATCAAGGCGAT 540  
Qy 543 TGGGGAAGTGAACCTGCTGTTTATGTCCTCAGAGAAATGCTTGGCTCTGAGCGAGAGAG 602  
Db 541 CGGGGAAGTGAACCTGCTGTTTATGTCCTCAGAGAAATGCTTGGCTCTGAGCGAGAGAG 600  
Qy 603 CTAGAGAAAGAGAACTGCTCTCTCTGCTTCTTAAAGAAAGAACTAAGATCCCTGAAATG 662  
Db 601 CTAGAGAAAGAGAACTGCTCTCTCTGCTTCTTAAAGAAAGAACTAAGATCCCTGAAATG 660  
Qy 663 GACTTTTTTACTAAGGAAAGTGAGAGCTAACGTCATCATCATTTAGAGAGATTTCAAT 722  
Db 661 GACTTTTTTACTAAGGAAAGTGAGAGCTAACGTCATCATCATTTAGAGATTTCAAT 720  
Qy 723 GAAACCTGGCTCAGTTGAAAAGAAATAGTGTCAAGTGTCCATGATGACACAGAGGTAGA 782  
Db 721 GAAACCTGGCTCAGTTGAAAAGAAATAGTGTCAAGTGTCCATGATGACACAGAGGTAGA 780

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QY 783 CTTGATACCAACAAAGTTCATTGACAAATATTTATTTGTCACTGATGATACAAACAGAAA 842
DB 781 CTTGATACCAACAAAGTTCATTGACAAATATTTATTTGTCACTGATGATACAAACAGAAA 840
QY 843 ATAATGTACTTTAAATAATTTTGAAGGAGGTACCTCTCATCTCTTTAGAAAAAAG 902
DB 841 AGTATGTACTTTAAATAATTTTGAAGGAGGTACCTCTCATCTCTTAGAAGAAAAG 900
QY 903 CTTATGTAACTTCATTTCATATCCAAATATTTATATATATGTAAGTTTATTTATAAGT 962
DB 901 CCTATGTAACTTCATTTCATATCCAAATATTTATATATATGTAAGTTTATTTATAAGT 960
QY 963 ATACATTTTATTTATGTACGTTTATTAATATGATTTATTTATAGAAAACATTTCTGCTA 1022
DB 961 ATACATTTTATTTATGTACGTTTATTAATATGATTTTATTTATAGAAAACATTTCTGATG 1020
QY 1023 TTGATATTT-AGTATAAGGCAATAATATT 1051
DB 1021 TTGATATTTGAGTATAAGCAATAATATT 1050

RESULT 11
US-10-807-837-10
; Sequence 10, Application US/10807837
; Publication No. US20040209330A1
; GENERAL INFORMATION:
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Chandraseker, Yasmin A.
; APPLICANT: Dillon, Stacey R.
; APPLICANT: Lehner, Joyce M.
; APPLICANT: Siadak, Anthony W.
; APPLICANT: Sivakumar, Pallavur V.
; APPLICANT: Moore, Margaret D.
; TITLE OF INVENTION: ANTI-IL-22RA ANTIBODIES AND METHODS OF USING IN INFLAMMATION
; FILE REFERENCE: 03-02
; CURRENT APPLICATION NUMBER: US/10/807,837
; CURRENT FILING DATE: 2004-03-24
; PRIOR APPLICATION NUMBER: US 60/457,481
; PRIOR FILING DATE: 2003-03-24
; PRIOR APPLICATION NUMBER: US 60/523,295
; PRIOR FILING DATE: 2003-11-17
; NUMBER OF SEQ ID NOS: 62
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 1050
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (5)...(589)
US-10-807-837-10

Query Match 88.8%; Score 993.2; DB 8; Length 1050;
Best Local Similarity 97.2%; Pred. No. 2.2e-197;
Matches 1021; Conservative 0; Mismatches 28; Indels 1; Gaps 1;

QY 3 AACAGGCTCTCCTCTCACTTATCAACTGTTTGACACTGTGGGATCTCTGATGCTGTCCT 62
DB 1 AACAGGCTCTCCTCTCACTTATCAACTTGTGCGATCGGTGATGCTGTCCT 60
QY 63 GCAGAAATCTATGAGTTTTCCTTATGGGACTTTTGGCCGCGACGCTGCTGCTTCAT 122
DB 61 GCAGAAATCTATGAGTTTTCCTTATGGGACTTTTGGCCGCGACGCTGCTGCTTCAT 120
QY 123 TGGCCTGTGGCCCGCAGGAGCAATATGCGCTGCCGTCACACCCGGTGCAAGCTTGAGGT 182
DB 121 TGGCCTGTGGCCCGCAGGAGCAATATGCGCTGCCCATCAACACCCGGTGCAAGCTTGAGGT 180
QY 183 GTCCAACTTCAGAGCCGCTACATCGTCAACCCGACCTTTATGCTGGCCAGGAGGCCAG 242
DB 181 GTCCAACTTCAGAGCCGCTACATCGTCAACCCGACCTTTATGCTGGCCAGGAGGCCAG 240
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QY 243 CTTTGAGATACAAACACAGAGCTCCGGCTCATCGGGAGAAAATGTTCCGAGGAGTCAG 302
DB 241 CTTTGAGATACAAACACAGAGCTCCGGCTCATCGGGAGAAAATGTTCCGAGGAGTCAG 300
QY 303 TGTAAAGATCAGTGTCTGATGAGCAGAGTGCTCAACTTCACCTCGAGAGCGTTCT 362
DB 301 TGTAAAGATCAGTGTCTGATGAGCAGAGTGCTCAACTTCACCTCGAGAGCATTTCT 360
QY 363 GCTCCCCCAGTCAGACAGGTTCCAGGCCCTACATGAGGAGGTGCTACCTTTCTCGACCA 422
DB 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATGAGGAGGTGCTGCTTTCTCGACCA 420
QY 423 ACTCAGCAATCAGCTCAGTGTCTGTCATCAGCGGTGACGACAGCAACATCCAGAGAA 482
DB 421 ACTCAGCAATCAGCTCAGTGTCTGTCATCAGTGTGACGACAGCAACATCCAGAGAA 480
QY 483 TGTCAAGAGCTCAGGAGGACAGTGAAGAGCTTGGAGAGTGGAGAGATCAAGCGAT 542
DB 481 TGTCAAGAGCTCAGGAGGACAGTGAAGAGCTTGGAGAGGCGGAGAGATCAAGCGAT 540
QY 543 TGGGGAACCTGGACCTGCTTTATGTCCTGAGAAAATGCTTGGCTCTGAGCGAGAGAAAG 602
DB 541 CGGGGAACCTGGACCTGCTTTATGTCCTGAGAAAATGCTTGGCTCTGAGCGAGAGAAAG 600
QY 603 CTAGAAAAAGAGAACTGCTTCTGCTCTTCTAAAGAAACAATAAGATCCCTGAATG 662
DB 601 CTAGAAAAAGAGAACTGCTTCTGCTCTTCTAAAGAAACAATAAGATCCCTGAATG 660
QY 663 GACTTTTTTACTAAAGGAAGTGAAGAGCTTAAGCTCCATCATCTTAGAGAGATTTCAAT 722
DB 661 GACTTTTTTACTAAAGGAAGTGAAGAGCTTAAGCTCCATCATCTTAGAGAGATTTCAAT 720
QY 723 GAAACCTGGCTCAGTTGAAAAAGAAAATAGTGTCAAGTGTCCATGAGACAGAGGTAGA 782
DB 721 GAAACCTGGCTCAGTTGAAAAAGAAAATAGTGTCAAGTGTCCATGAGACAGAGGTAGA 780
QY 783 CTTGATAACCAACAAAGATTCATTGACAAATATTTATTTGTCACTGATGATACAAACAGAAA 842
DB 781 CTTGATAACCAACAAAGATTCATTGACAAATATTTATTTGTCACTGATGATACAAACAGAAA 840
QY 843 ATAATGTACTTTAAAAAATTTTGAAGGAGGTACCTCTCATCTTTAGAAAAAAG 902
DB 841 AGTATGTACTTTAAAAAATTTTGAAGGAGGTACCTCTCATCTTAGAAGAAAAG 900
QY 903 CTTATGTAACTTCATTTCATATCCAAATATTTATATATATGTAAGTTTATTTATAAGT 962
DB 901 CCTATGTAACTTCATTTCATATCCAAATATTTATATATGTAAGTTTATTTATAAGT 960
QY 963 ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAAACATTTCTGCTA 1022
DB 961 ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAAACATTTCTGATG 1020
QY 1023 TTGATATTT-AGTATAAGGCAATAATATT 1051
DB 1021 TTGATATTTGAGTATAAGCAATAATATT 1050
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RESULT 12
US-10-968-432-42
; Sequence 42, Application US/10968432
; Publication No. US20050065321A1
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Chen, Zhi
; APPLICANT: Hughes, Steven D.
; TITLE OF INVENTION: Human Cytokine Receptor
; FILE REFERENCE: 01-12C1
; CURRENT APPLICATION NUMBER: US/10/968,432
; CURRENT FILING DATE: 2004-10-19
; PRIOR APPLICATION NUMBER: US 60/279,222
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; PRIOR FILING DATE: 2001-03-27
; PRIOR APPLICATION NUMBER: US 10/104,919
; PRIOR FILING DATE: 2002-03-22
; NUMBER OF SEQ ID NOS: 62
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 42
; LENGTH: 1050
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (5)...(589)
US-10-968-432-42

Query Match      88.8%; Score 993.2; DB 9; Length 1050;
Best Local Similarity 97.2%; Pred. No. 2.2e-197;
Matches 1021; Conservative 0; Mismatches 28; Indels 1; Gaps 1;

Qy  3 AACAGGCTCTCTCTCACTTATCAACTTGTGACACTTGTGCGATCTCTGATGGCTGTCCT 62
Db  1 AACAGGCTCTCTCTCACTTATCAACTTGTGACACTTGTGCGATCGGTGATGGCTGTCCT 60

Qy  63 GCAGAAATCTATGATGTTTTTCCCTTATGGGACTTTGGCCGACGCTGCCCTGCTTCTCAT 122
Db  61 GCAGAAATCTATGATGTTTTTCCCTTATGGGACTTTGGCCGACGCTGCCCTGCTTCTCAT 120

Qy  123 TGCCTGTGGCCGACGAGGCAATGCGTCCGTCNAACCCGCGTGAAGCTTGAGGT 182
Db  121 TGCCTGTGGCCGACGAGGCAATGCGTCCGTCNAACCCGCGTGAAGCTTGAGGT 180

Qy  183 GTCCAACTTCCAGCAGCGGTACATCGTCAACCGCACTTTATGCTGGCCAGAGGCCAG 242
Db  181 GTCCAACTTCCAGCAGCGGTACATCGTCAACCGCACTTTATGCTGGCCAGAGGCCAG 240

Qy  243 CCTTGCAGATAACCAACACAGACGTCGCGCTCATCGGGAGAAACTGTTCCGAGGAGTCAG 302
Db  241 CCTTGCAGATAACCAACACAGACGTCGCGCTCATCGGGAGAAACTGTTCCGAGGAGTCAG 300

Qy  303 TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACTCCCTGGAGAGCTTCT 362
Db  301 TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACTCCCTGGAGAGATCTCT 360

Qy  363 GCTCCCCAGTCAGACAGGTTCCAGCCCTCATATGCGAGGTTGCTTCTTCCCTGACCAA 422
Db  361 GCTCCCCAGTCAGACAGGTTCCGGCCCTCATATGCGAGGTTGCTTCTTCCCTGACCAA 420

Qy  423 ACTCAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGAACGACGAGCAATCCAGAGAA 482
Db  421 ACTCAGCAATCAGCTCAGCTCCTGTCAATCAGTGGTGAACGACGAGCAATCCAGAGAA 480

Qy  483 TGTCAAGAGGCTGAAGGACAGTGAAGAAAGCTTTGGAGAGTGGAGAGATCAAGGCCAT 542
Db  481 TGTCAAGAGGCTGAAGGACAGTGAAGAAAGCTTTGGAGAGTGGAGAGATCAAGGCCAT 540

Qy  543 TGGGGAACCTGACCTGCTGTTTATGCTCTCAGAGAAAGCTTGTGCGTCTGAGCGAGAGAG 602
Db  541 CGGGGAACCTGACCTGCTGTTTATGCTCTCAGAGAAAGCTTGTGCGTCTGAGCGAGAGAG 600

Qy  603 CTAGAAAACGAGAACTGCTCTCTTCTGCTTCTTAAAGAAACAAATAGATCCCTGAATG 662
Db  601 CTAGAAAACGAGAACTGCTCTCTTCTGCTTCTTAAAGAAACAAATAGATCCCTGAATG 660

Qy  663 GACTTTTTTACTAAGGAAGTGAAGACTAAGCTTCAATCATCATATTAGAAATTTTCAT 722
Db  661 GACTTTTTTACTAAGGAAGTGAAGACTAAGCTTCAATCATCATATTAGAAATTTTCAT 720

Qy  723 GAAACCTGGCTCAGTTGAAAAGAAAATAGTGTCAAAGTTGTCCATGACACAGAGGTAGA 782
Db  721 GAAACCTGGCTCAGTTGAAAAGAAAATAGTGTCAAAGTTGTCCATGACACAGAGGTAGA 780

Qy  783 CTTGATAACCAAGATTTCAATTTGACAAATATTTTATTTGTCATCTGATGATACAAACAGAAA 842
Db  781 CTTGATAACCAAGATTTCAATTTGACAAATATTTTATTTGTCATTTGATGATGCAACAGAAA 840

; PRIOR FILING DATE: 2001-03-27
; PRIOR APPLICATION NUMBER: US 11/045,944
; PRIOR FILING DATE: 2005-01-28
; CURRENT FILING DATE: 2005-01-28
; PRIOR APPLICATION NUMBER: US 60/273,035
; PRIOR FILING DATE: 2001-03-02
; PRIOR APPLICATION NUMBER: US 60/279,232
; PRIOR FILING DATE: 2001-03-27
; PRIOR APPLICATION NUMBER: US 10/090,365
; PRIOR FILING DATE: 2002-03-04
; NUMBER OF SEQ ID NOS: 49
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 40
; LENGTH: 1050
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (50)...(589)
US-11-045-944-40

RESULT 13
US-11-045-944-40
; Sequence 40, Application US/11045944
; Publication No. US20050124796A1
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Chen, Zhi
; TITLE OF INVENTION: Mouse Cytokine Receptor
; FILE REFERENCE: 01-08D1
; CURRENT APPLICATION NUMBER: US/11/045,944
; CURRENT FILING DATE: 2005-01-28
; PRIOR APPLICATION NUMBER: US 60/273,035
; PRIOR FILING DATE: 2001-03-02
; PRIOR APPLICATION NUMBER: US 60/279,232
; PRIOR FILING DATE: 2001-03-27
; PRIOR APPLICATION NUMBER: US 10/090,365
; PRIOR FILING DATE: 2002-03-04
; NUMBER OF SEQ ID NOS: 49
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 40
; LENGTH: 1050
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (50)...(589)
US-11-045-944-40

Query Match      88.8%; Score 993.2; DB 10; Length 1050;
Best Local Similarity 97.2%; Pred. No. 2.2e-197;
Matches 1021; Conservative 0; Mismatches 28; Indels 1; Gaps 1;

Qy  3 AACAGGCTCTCTCTCACTTATCAACTTGTGACACTTGTGCGATCTCTGATGGCTGTCCT 62
Db  1 AACAGGCTCTCTCTCACTTATCAACTTGTGACACTTGTGCGATCTCTGATGGCTGTCCT 60

Qy  63 GCAGAAATCTATGATGTTTTTCCCTTATGGGACTTTGGCCGACGCTGCCCTGCTTCTCAT 122
Db  61 GCAGAAATCTATGATGTTTTTCCCTTATGGGACTTTGGCCGACGCTGCCCTGCTTCTCAT 120

Qy  123 TGCCTGTGGCCGACGAGGCAATGCGTCCGTCNAACCCGCGTGAAGCTTGAGGT 182
Db  121 TGCCTGTGGCCGACGAGGCAATGCGTCCGTCNAACCCGCGTGAAGCTTGAGGT 180

Qy  183 GTCCAACTTCCAGCAGCGGTACATCGTCAACCGCACTTTATGCTGGCCAGAGGCCAG 242
Db  181 GTCCAACTTCCAGCAGCGGTACATCGTCAACCGCACTTTATGCTGGCCAGAGGCCAG 240

Qy  243 CCTTGCAGATAACCAACACAGACGTCGCGCTCATCGGGAGAAACTGTTCCGAGGAGTCAG 302
Db  241 CCTTGCAGATAACCAACACAGACGTCGCGCTCATCGGGAGAAACTGTTCCGAGGAGTCAG 300

Qy  303 TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACTCCCTGGAGAGCTTCT 362
Db  301 TGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACTCCCTGGAGAGATCTCT 360

Qy  363 GCTCCCCAGTCAGACAGGTTCCAGCCCTCATATGCGAGGTTGCTTCTTCCCTGACCAA 422
Db  361 GCTCCCCAGTCAGACAGGTTCCGGCCCTCATATGCGAGGTTGCTTCTTCCCTGACCAA 420

Qy  423 ACTCAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGAACGACGAGCAATCCAGAGAA 482
Db  421 ACTCAGCAATCAGCTCAGCTCCTGTCAATCAGTGGTGAACGACGAGCAATCCAGAGAA 480

Qy  483 TGTCAAGAGGCTGAAGGACAGTGAAGAAAGCTTTGGAGAGTGGAGAGATCAAGGCCAT 542
Db  481 TGTCAAGAGGCTGAAGGACAGTGAAGAAAGCTTTGGAGAGTGGAGAGATCAAGGCCAT 540

Qy  543 TGGGGAACCTGACCTGCTGTTTATGCTCTCAGAGAAAGCTTGTGCGTCTGAGCGAGAGAG 602
Db  541 CGGGGAACCTGACCTGCTGTTTATGCTCTCAGAGAAAGCTTGTGCGTCTGAGCGAGAGAG 600

Qy  603 CTAGAAAACGAGAACTGCTCTCTTCTGCTTCTTAAAGAAACAAATAGATCCCTGAATG 662
Db  601 CTAGAAAACGAGAACTGCTCTCTTCTGCTTCTTAAAGAAACAAATAGATCCCTGAATG 660

Qy  663 GACTTTTTTACTAAGGAAGTGAAGACTAAGCTTCAATCATCATATTAGAAATTTTCAT 722
Db  661 GACTTTTTTACTAAGGAAGTGAAGACTAAGCTTCAATCATCATATTAGAAATTTTCAT 720

Qy  723 GAAACCTGGCTCAGTTGAAAAGAAAATAGTGTCAAAGTTGTCCATGACACAGAGGTAGA 782
Db  721 GAAACCTGGCTCAGTTGAAAAGAAAATAGTGTCAAAGTTGTCCATGACACAGAGGTAGA 780

Qy  783 CTTGATAACCAAGATTTCAATTTGACAAATATTTTATTTGTCATCTGATGATACAAACAGAAA 842
Db  781 CTTGATAACCAAGATTTCAATTTGACAAATATTTTATTTGTCATTTGATGATGCAACAGAAA 840
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Db 301 TGTCTAAGGATCAGTGTCTACCTGATGAGCAGGCTGCTCAACTTCCACCTCGAAGACATTTCT 360
Qy 363 GCTCCCCCAGCTCAGACAGGTTTCAGCCCTACATGTCAGGAGTGATACCTTTCTCTGACCAA 422
Db 361 GCTCCCCCAGCTCAGACAGGTTTCAGCCCTACATGTCAGGAGTGATGCTTTCTCTGACCAA 420
Qy 423 ACTCAGCAATCAGCTCAGCTCCTGTGTCTCATCAGCGGTGACGACAGCAACAATCCAGAAGAA 482
Db 421 ACTCAGCAATCAGCTCAGCTCCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 480
Qy 483 TGTCTAAGGCTCAGGAGACAGTGAAGAGCTTGGAGAGTGGAGAGATCAAGCGAT 542
Db 481 TGTCTAAGGCTCAGGAGACAGTGAAGAGCTTGGAGAGTGGAGAGATCAAGCGAT 540
Qy 543 TGTCTAAGGCTCAGGAGACAGTGAAGAGCTTGGAGAGTGGAGAGATCAAGCGAT 602
Db 541 CGGGGAACCTGACCTGCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 600
Qy 603 CTAGAAAACGAGAACTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 662
Db 601 CTAGAAAACGAGAACTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 660
Qy 663 GACTTTTTTACTAAGGAAAGTGAAGAGCTTAAAGCTTAAAGCTTAAAGCTTAAAGCTTAAAG 722
Db 661 GACTTTTTTACTAAGGAAAGTGAAGAGCTTAAAGCTTAAAGCTTAAAGCTTAAAGCTTAAAG 720
Qy 723 GAAACCTGGCTCAGTGTGAAAAGAAATAGTGTCAAGTGTGTGTGTGTGTGTGTGTGTGTGTGT 782
Db 721 GAAACCTGGCTCAGTGTGAAAAGAAATAGTGTCAAGTGTGTGTGTGTGTGTGTGTGTGTGTGT 780
Qy 783 CTTGATAACCAAGAAATCAATTGACAAATATTTATTTATTTATTTATTTATTTATTTATTTATTT 842
Db 781 CTTGATAACCAAGAAATCAATTGACAAATATTTATTTATTTATTTATTTATTTATTTATTTATTT 840
Qy 843 ATATGTACTTTTAAAGAAATTTTGAAGAGGTTACCTCTCATTTAGAAAAG 902
Db 841 AGTATGTACTTTTAAAGAAATTTTGAAGAGGTTACCTCTCATTTAGAAAAG 900
Qy 903 CTTATGTACTTTTAAAGAAATTTTGAAGAGGTTACCTCTCATTTAGAAAAG 962
Db 901 CTTATGTACTTTTAAAGAAATTTTGAAGAGGTTACCTCTCATTTAGAAAAG 960
Qy 963 ATACATTTTATTTATGTGAGTTTATTAATATGAGTTTATTTATAGAAAACATTTATCTGCTA 1022
Db 961 ATACATTTTATTTATGTGAGTTTATTAATATGAGTTTATTTATAGAAAATTTATCTGATG 1020
Qy 1023 TTGATATTT-AGTATAGGCAATAATATT 1051
Db 1021 TTGATATTTGAGTATAGGCAATAATATT 1050
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## RESULT 14

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US-09-746-375-37
; Sequence 37, Application US/09746375
; Publication No. US20030170823A1
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
; APPLICANT: Kindsvogel, Wayne
; TITLE OF INVENTION: NOVEL CYTOKINE ZCYTO18
; FILE REFERENCE: 99-106
; CURRENT APPLICATION NUMBER: US/09/746,375
; CURRENT FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 60/172,105
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: US 60/****,***
; PRIOR FILING DATE: 2000-12-01
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 37
; LENGTH: 778
; TYPE: DNA
; ORGANISM: mus musculus
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; FEATURE:
; NAME/KEY: CDS
; LOCATION: (47) ... (583)
US-09-746-375-37

Query Match 68.7%; Score 768.4; DB 3; Length 778;
Best Local Similarity 99.8%; Pred. No. 1.9e-150;
Matches 772; Conservative 0; Mismatches 6; Indels 0; Gaps 0;

Qy 6 AGGCTCTCTCTCCTCACTTATCAACTGTGTGACACACTGTGTGCGGATCTCTGATGGCTGTCTCTGCA 65
Db 1 AGGCTCTCTCTCCTCACTTATCAACTGTGTGACACTGTGTGCGGATCGGTGATGCTGTCTCTGCA 60
Qy 66 GAAATCTATGAGTTTTTCCCTTATGGGACTTTTGGCGCCAGCTGCTCTCTCTCTCTCTCTCTCTCT 125
Db 61 GAAATCTATGAGTTTTTCCCTTATGGGACTTTTGGCGCCAGCTGCTCTCTCTCTCTCTCTCTCTCT 120
Qy 126 CTTGTGGGCCCAGGAGCAAAATGCGTGCCTGCTCAACACCCGGTGCAGCTTGAAGGTGTC 185
Db 121 CTTGTGGGCCCAGGAGCAAAATGCGTGCCTGCTCAACACCCGGTGCAGCTTGAAGGTGTC 180
Qy 186 CAACTTCCAGCAGCGCTAGCATCGTCAACCGCACCTTTATGCTGGCCAAAGGAGGCCAGCCT 245
Db 181 CAACTTCCAGCAGCGCTAGCATCGTCAACCGCACCTTTATGCTGGCCAAAGGAGGCCAGCCT 240
Qy 246 TGCAGATAACAACAACAAGCTCCGGCTCATCGGGGAGAAACTGTTCGAGGAGTCACTGTC 305
Db 241 TGCAGATAACAACAACAAGCTCCGGCTCATCGGGGAGAAACTGTTCGAGGAGTCACTGTC 300
Qy 306 TAAAGATCAGTGTCTACTGATGAAGCAGGTGCTCAACTTCACTTCACTTCACTTCACTTCACTT 365
Db 301 TAAAGATCAGTGTCTACTGATGAAGCAGGTGCTCAACTTCACTTCACTTCACTTCACTTCACTT 360
Qy 366 CCCCAGTCAAGACAGGTTCCAGCCCTACATGTCAGGAGGTGCTACCTTTCTCTGACCAAACT 425
Db 361 CCCCAGTCAAGACAGGTTCCAGCCCTACATGTCAGGAGGTGCTGCTTTCTCTGACCAAACT 420
Qy 426 CAGCAATCAGCTCAGTCTCTGTCTCATCATCAGCGGTGACGACAGAAACATCCAGAAAGATGT 485
Db 421 CAGCAATCAGCTCAGTCTCTGTCTCATCATCAGCGGTGACGACAGAAACATCCAGAAAGATGT 480
Qy 486 CAGAGGCTGAGGAGACAGTGAAGAGCTTGGAGAGTGGAGAGATCAAGCGATTGG 545
Db 481 CAGAGGCTGAGGAGACAGTGAAGAGCTTGGAGAGTGGAGAGATCAAGCGATTGG 540
Qy 546 GGAACCTGGACCTCTCTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTA 605
Db 541 GGAACCTGGACCTCTCTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTA 600
Qy 606 GAAACCGAAGAACTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 665
Db 601 GAAACCGAAGAACTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 660
Qy 666 TTTTCTTAAAGAAAGTGAAGCTAAGCTCATCATCATCTATTAGAGATTTCATGAA 725
Db 661 TTTTCTTAAAGAAAGTGAAGCTAAGCTCATCATCATCTATTAGAGATTTCATGAA 720
Qy 726 ACCTGGCTCAGTTGAAAAAGAAATAGTGTCAAGTTGTCTCATGAGACGAGAGGTAGAC 783
Db 721 ACCTGGCTCAGTTGAAAAAGAAATAGTGTCAAGTTGTCTCATGAGACGAGAGGTAGAC 778
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## RESULT 15

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US-10-395-741B-37
; Sequence 37, Application US/10395741B
; Publication No. US20040023341A1
; GENERAL INFORMATION:
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Hughes, Steven D.
; APPLICANT: Chandrasekhar, Yasmin A.
; TITLE OF INVENTION: ANTI-IL-17F ANTIBODIES AND METHODS OF
; TITLE OF INVENTION: USING IN INFLAMMATION
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; FILE REFERENCE: 02-04  
; CURRENT APPLICATION NUMBER: US/10/395,741B  
; CURRENT FILING DATE: 2003-03-24  
; PRIOR APPLICATION NUMBER: US 60/366,842  
; PRIOR FILING DATE: 2002-03-22  
; NUMBER OF SEQ ID NOS: 67  
; SOFTWARE: FastSeq for Windows Version 3.0  
; SEQ ID NO 37  
; LENGTH: 778  
; TYPE: DNA  
; ORGANISM: mus musculus  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (47)...(583)  
US-10-395-741B-37

Query Match 68.7%; Score 768.4; DB 7; Length 778;  
Best Local Similarity 99.2%; Pred. No. 1.9e-150;  
Matches 772; Conservative 0; Mismatches 6; Indels 0; Gaps 0;  
  
Qy 6 AGGCTCTCTCTCACTTATCAACTTGTGACACACTTGTGGGATCTCTGATGGCTGTCTCTGCA 65  
Db |||||||  
1 AGGCTCTCTCTCACTTATCAACTTGTGACACACTTGTGCGATCGGTGATGGCTGTCTCTGCA 60  
Qy 66 GAAATCTATGAGTTTTCCTTTATGGGACTTTGCGCCGACAGTGCCTTCTCTCATTTGC 125  
Db |||||||  
61 GAAATCTATGAGTTTTCCTTTATGGGACTTTGCGCCGACAGTGCCTTCTCTCATTTGC 120  
Qy 126 CCTGTGGGCCCAGGAGGCAATGGCGTGCCTGCAACACCCGGTGCAGCTTTGAGGTGTC 185  
Db |||||||  
121 CCTGTGGGCCCAGGAGGCAATGGCGTGCCTGCAACACCCGGTGCAGCTTTGAGGTGTC 180  
Qy 186 CAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGTGGCCAAAGGAGGCGAGCCT 245  
Db |||||||  
181 CAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGTGGCCAAAGGAGGCGAGCCT 240  
Qy 246 TGCAGATAACACACAGACGTCGGCTCATCGGGAGAACTGTTCCGAGGAGTCAGTGC 305  
Db |||||||  
241 TGCAGATAACACACAGATGTCGGCTCATCGGGAGAACTGTTCCGAGGAGTCAGTGC 300  
Qy 306 TAAAGATCAGTGTACCTGTAGTGAAGCAGGTGCTCAACTTACCTCTGGAAGACGTTCTGCT 365  
Db |||||||  
301 TAAAGATCAGTGTACCTGTAGTGAAGCAGGTGCTCAACTTACCTCTGGAAGACGTTCTGCT 360  
Qy 366 CCCCAGTCAACACAGGTTCCAGCCCTACATGAGGAGGTGTA CTTTCTTGACCAAACT 425  
Db |||||||  
361 CCCCAGTCAACACAGGTTCCAGCCCTACATGAGGAGGTGTA CTTTCTTGACCAAACT 420  
Qy 426 CAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGACGACCAAGAACATCCAGAGAAATGT 485  
Db |||||||  
421 CAGCAATCAGCTCAGCTCCTGTCAATCAGCGGTGACGACCAAGAACATCCAGAGAAATGT 480  
Qy 486 CAGAAGGCTGAAGGAGACAGTGAAAGCTTTGGAGAGAGTGGAGAGATCAAGGGGATTGG 545  
Db |||||||  
481 CAGAAGGCTGAAGGAGACAGTGAAAGCTTTGGAGAGAGTGGAGAGATCAAGGGGATTGG 540  
Qy 546 GGAATCGAACCTGTGTTTATGTCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTA 605  
Db |||||||  
541 GGAATCGAACCTGTGTTTATGTCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTA 600  
Qy 606 GAAACGAGAACTGCTCCTCTCCCTTCTAAAGAAACAATAGATCCCTGATGGAC 665  
Db |||||||  
601 GAAACGAGAACTGCTCCTCTCCCTTCTAAAGAAACAATAGATCCCTGATGGAC 660  
Qy 666 TTTTCTTAAAGGAAAGTGAGAACTAACCGTCCATCATTTAGAGATTTCACATGAA 725  
Db |||||||  
661 TTTTCTTAAAGGAAAGTGAGAACTAACCGTCCATCATTTAGAGATTTCACATGAA 720  
Qy 726 ACCTGGCTCAGTTGAAAAAGAAAAATAGTGTCAAGTTGTCATGAGACCAGAGGTAGAC 783  
Db |||||||  
721 ACCTGGCTCAGTTGAAAAAGAAAAATAGTGTCAAGTTGTCATGAGACCAGAGGTAGAC 778

Search completed: December 21, 2005, 21:07:05  
Job time : 562.633 secs



GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 78.0229 Seconds  
(without alignments)  
7442.822 Million cell updates/sec

Title: US-09-751-797-7  
Perfect score: 1119  
Sequence: 1 taacaggctctctctcac.....tggatatcataaaaaaaa 1119

Scoring table: IDENTITY NUC  
Gapop 10.0 , Gapext 1.0

Searched: 4168288 seqs, 259477437 residues

Total number of hits satisfying chosen parameters: 8336576

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Published Applications NA.New.\*  
1: /cgn2\_6/ptodata/2/pubpna/US08\_NEW\_PUB.seq.\*  
2: /cgn2\_6/ptodata/2/pubpna/US06\_NEW\_PUB.seq.\*  
3: /cgn2\_6/ptodata/2/pubpna/US07\_NEW\_PUB.seq.\*  
4: /cgn2\_6/ptodata/2/pubpna/PCT\_NEW\_PUB.seq.\*  
5: /cgn2\_6/ptodata/2/pubpna/US09\_NEW\_PUB.seq.\*  
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8: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq2.\*  
9: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq3.\*  
10: /cgn2\_6/ptodata/2/pubpna/US60\_NEW\_PUB.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	1119	100.0	1119	7	US-11-177-987-7
2	1047.8	93.6	1111	7	US-11-177-987-9
3	601.4	53.7	7445	7	US-11-177-987-8
4	555.2	49.6	5935	7	US-11-177-987-42
5	524.8	46.9	1152	7	US-11-102-240-153
6	409.2	36.6	690	7	US-11-177-987-25
7	223.6	20.0	418	7	US-11-177-987-18
8	126	11.3	4797	7	US-11-177-987-26
9	54.6	4.9	26772	6	US-10-995-561-13313
10	54.6	4.9	54946	6	US-10-995-561-13479
11	54	4.8	171486	7	US-11-121-086-105
12	53.8	4.8	130276	6	US-10-661-966-1
13	53.6	4.8	49979	6	US-10-995-561-13443
14	53.4	4.8	47444	6	US-10-995-561-13354
15	52.8	4.7	173602	7	US-11-121-086-25
16	52.6	4.7	1819	6	US-10-750-185-29496
17	52.4	4.7	184868	7	US-11-121-086-88
18	52.2	4.7	201990	6	US-10-995-561-13303
19	51.8	4.6	173602	7	US-11-121-086-25
20	51.2	4.6	161874	7	US-11-121-086-75
21	51	4.6	119036	6	US-10-995-561-13314
22	51	4.6	151169	7	US-11-121-086-38
23	50.6	4.5	151169	7	US-11-121-086-38

ALIGNMENTS

RESULT 1

US-11-177-987-7  
; Sequence 7, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT FILING DATE: 2005-07-07  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; PRIOR FILING DATE: 2005-07-07  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-11-177-987-7  
Query Match 100.0%; Score 1119; DB 7; Length 1119;  
Best Local Similarity 100.0%; Pred. No. 1.6e-245;  
Matches 1119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 1 TAAACAGGCTCTCTCTCACTTATCACTTGTGACACTTGTGACATCTGTGATGCGTCTC 60  
Db 1 TAAACAGGCTCTCTCTCACTTATCACTTGTGACACTTGTGACATCTGTGATGCGTCTC 60  
Qy 61 CTCGAGAAATCTATGAGTTTCCCTTATGCGGACTTTGCGGACCTGCTGCTTCTC 120  
Db 61 CTCGAGAAATCTATGAGTTTCCCTTATGCGGACTTTGCGGACCTGCTGCTTCTC 120  
Qy 121 ATTGCCCCGTGGGCCAGGAGGCAAAATGCGCTGCCGTCAACACCCGGTGCAAGCTTGAG 180  
Db 121 ATTGCCCCGTGGGCCAGGAGGCAAAATGCGCTGCCGTCAACACCCGGTGCAAGCTTGAG 180  
Qy 181 GTGTCAACTTCCAGCAGCCGTTACATGTCACCGACCTTTATGCTGTGGCCAGGAGGCC 240  
Db 181 GTGTCAACTTCCAGCAGCCGTTACATGTCACCGACCTTTATGCTGTGGCCAGGAGGCC 240

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Db 181 GTGTCCAATTCACAGCGCGTACATCGTCAACCGCACCTTTATGTGCGCAAGAGGCC 240
Qy 241 AGCCTTCAGATACACACAGAGCTCGGCTCATCGGGAGAACTGTTCCGAGGATC 300
Db 241 AGCCTTCAGATACACACAGAGCTCGGCTCATCGGGAGAACTGTTCCGAGGATC 300
Qy 301 AGTGTAAAGATCAGTCTACCTCATGAAGCAGGTGCTCAACTTCACCTCGGAAGAGCTT 360
Db 301 AGTGTAAAGATCAGTCTACCTCATGAAGCAGGTGCTCAACTTCACCTCGGAAGAGCTT 360
Qy 361 CTGCTCCCCCAGTCAGACAGGTTCACGCCCTPACATGACAGAGGTGTAACCTTCTCAGC 420
Db 361 CTGCTCCCCCAGTCAGACAGGTTCACGCCCTPACATGACAGAGGTGTAACCTTCTCAGC 420
Qy 421 AAATCAGCAATCAGCTCAGCTCTGTGCATCATCAGCGGTGACGACCAAGATCCAGAAG 480
Db 421 AAATCAGCAATCAGCTCAGCTCTGTGCATCATCAGCGGTGACGACCAAGATCCAGAAG 480
Qy 481 AATGTCAAGAGGCTGAAGGAGACAGTGAAGAGCTTGGAGAGCTGGAGAGATCAAGGCG 540
Db 481 AATGTCAAGAGGCTGAAGGAGACAGTGAAGAGCTTGGAGAGCTGGAGAGATCAAGGCG 540
Qy 541 ATTGGGAACTGGACCTGCTGCTTTATGTCTCTGAGAAATGCTTGCCTCTGAGCGAGAAGA 600
Db 541 ATTGGGAACTGGACCTGCTGCTTTATGTCTCTGAGAAATGCTTGCCTCTGAGCGAGAAGA 600
Qy 601 AGCTAGAAAACGAAGAACTGCTCTTCTGCTTCTTAAAGAAACAATAGATCCCTGAA 660
Db 601 AGCTAGAAAACGAAGAACTGCTCTTCTGCTTCTTAAAGAAACAATAGATCCCTGAA 660
Qy 661 TGGACTTTTTTACTAAAGAAAGTGAAGGCTAAGCTCATCATCTATTAGAAAGATTTTAC 720
Db 661 TGGACTTTTTTACTAAAGAAAGTGAAGGCTAAGCTCATCATCTATTAGAAAGATTTTAC 720
Qy 721 ATGAAACCTGGCTCAGTTGAAAAAGAAATAGTGTCAAGTTGTCCATGAGACCAAGGTA 780
Db 721 ATGAAACCTGGCTCAGTTGAAAAAGAAATAGTGTCAAGTTGTCCATGAGACCAAGGTA 780
Qy 781 GACTGTATACCAAGATTCATGTACATATTTTATTTATTTATGTCACTGATGATACACAGAA 840
Db 781 GACTGTATACCAAGATTCATGTACATATTTTATTTATTTATGTCACTGATGATACACAGAA 840
Qy 841 AAATAAGTACTTTTAAATAATTTTGAAGAGGTTACCTCTCATCTCTTTAGAAAAAA 900
Db 841 AAATAAGTACTTTTAAATAATTTTGAAGAGGTTACCTCTCATCTCTTTAGAAAAAA 900
Qy 901 AGCTTATGTAACCTTCAATTCATATCCATATTTTATATATATGTAAGTTTATTTATATA 960
Db 901 AGCTTATGTAACCTTCAATTCATATCCATATTTTATATATATGTAAGTTTATTTATATA 960
Qy 961 GTATACATTTTATTTATGTAGTTTATTAATATGATTTTATTTATAGAAACATTTCTGC 1020
Db 961 GTATACATTTTATTTATGTAGTTTATTAATATGATTTTATTTATAGAAACATTTCTGC 1020
Qy 1021 TATTGATATTTAGTATAGGCAATAATATTTATGACATAACTATGAAAAACAGATATC 1080
Db 1021 TATTGATATTTAGTATAGGCAATAATATTTATGACATAACTATGAAAAACAGATATC 1080
Qy 1081 TTAGGCTTTAATAACACATGGATATCATATAAAAAAAA 1119
Db 1081 TTAGGCTTTAATAACACATGGATATCATATAAAAAAAA 1119
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RESULT 2
US-11-177-987-9
; Sequence 9, Application US/11177987
; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
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; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 9
; LENGTH: 1111
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; US-11-177-987-9

Query Match 93.6%; Score 1047.8; DB 7; Length 1111;
Best Local Similarity 97.0%; Pred. No. 2.5e-229;
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;

Qy 3 AACAGGCTCTCTCTCACTTATCAACTGTGTGACACTTGTGGGATCTCTGATGGCTGTCT 62
Db 1 AACAGGCTCTCTCTCACTTATCAACTGTGTGACACTTGTGGGATCTCTGATGGCTGTCT 60
Qy 63 GCAGAAATCTATGAGTTTTCCTTATGGGACTTTTGGCCGCGCAGCTGCCCTTCTCAT 122
Db 61 GCAGAAATCTATGAGTTTTCCTTATGGGACTTTTGGCCGCGCAGCTGCCCTTCTCAT 120
Qy 123 TGCCTCTGGGCCCAGGAGGCAATGCGCTGCCCGTCAACACCCGGTGCAGCTTGAGGT 182
Db 121 TGCCTCTGGGCCCAGGAGGCAATGCGCTGCCCGTCAACACCCGGTGCAGCTTGAGGT 180
Qy 183 GTCCAACTTCAGCAGCGCTACATCGTCAACCGCACCTTTATGTGGCCAAAGAGGCCAG 242
Db 181 GTCCAACTTCAGCAGCGCTACATCGTCAACCGCACCTTTATGTGGCCAAAGAGGCCAG 240
Qy 243 CCTTGACAGTAAACACACAGACGCTCCGCTCATCGGGAGAGAACTGTTCGAGGAGTCT 302
Db 241 CCTTGACAGTAAACACACAGACGCTCCGCTCATCGGGAGAGAACTGTTCGAGGAGTCT 300
Qy 303 TGCTTAAAGATCAGTGTCTACCTGTATGAAGCAGGTGCTCAACTTTCACCTTGGAGAGCTTCT 362
Db 301 TGCTTAAAGATCAGTGTCTACCTGTATGAAGCAGGTGCTCAACTTTCACCTTGGAGAGATTTCT 360
Qy 363 GCTCCCCAGTCAAGAGGTTTCAGGCCCTACATCGAGAGGTGTACTCTTCTCGACCAA 422
Db 361 GCTCCCCAGTCAAGAGGTTTCAGGCCCTACATCGAGAGGTGTGTCTTCTCGACCAA 420
Qy 423 ACTCAGCAATCAGCTCAGCTCTCTGTCATCATCAGCGGTGACGACCAAGATCCAGAGAA 482
Db 421 ACTCAGCAATCAGCTCAGCTCTCTGTCATCATGAGTGGTGACGCCAAGATCCAGAGAA 480
Qy 483 TGTCAAGAGGCTGAAGGAGACAGTGAAGAAAGCTTTGGAGAGAGTGGAGAGATCAAGGCGAT 542
Db 481 TGTCAAGAGGCTGAAGGAGACAGTGAAGAAAGCTTTGGAGAGAGTGGAGAGATCAAGGCGAT 540
Qy 543 TGGGAACTGGACCTGCTGTTTATGTCTCTGAGAAATGCTTTCGCTCTGAGCGAGAGAG 602
Db 541 CGGGAACTGGACCTGCTGTTTATGTCTCTGAGAAATGCTTTCGCTCTGAGCGAGAGAG 600
Qy 603 CTAGAAACGAGAACTGCTCTCTTCTGCTTCTTAAAGAAACAATAGATCCCTGAAATG 662
Db 601 CTAGAAACGAGAACTGCTCTCTTCTGCTTCTTAAAGAAACAATAGATCCCTGAAATG 660
Qy 663 GACTTTTTTACTTAAAGGAAAGTGAAGAGCTTAACGTCATCATCATTTAGAAGATTTTACAT 722
Db 661 GACTTTTTTACTTAAAGGAAAGTGAAGAGCTTAACGTCATCATCATTTAGAAGATTTTACAT 720
Qy 723 GAAACCTGGCTCAGTTGAAAAAGAAATAGTGTCAAGTTGTCCATGAGACCAAGGATGA 782
Db 721 GAAACCTGGCTCAGTTGAAAAAGAAATAGTGTCAAGTTGTCCATGAGACCAAGGATGA 780
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Qy	690	GCTAACGTCCTCATCATATTAGAAATTTTCATGAAACCTGGCTCAGTTGAAAAGAAA	749
Db	5401	GCTAACGTCACCACTCATTTAGAAATTTTCATGAAACCTGGCTCAGTTGAAAAGAAA	5460
Qy	750	TAGTGTCAAGTTGTCCATGAGACGAGGTAGACTTGATAACCAAAAGATTTCATTGACA	809
Db	5461	TAGTGTCAAGTTGTCCATGAGACGAGGTAGACTTGATAACCAAAAGATTTCATTGACA	5520
Qy	810	ATATTTTATTGTCTCTGATGATACAACAGAAAAATATGTAATTTAAAAATTTGTTGAA	869
Db	5521	ATATTTTATTGTCTCTGATGATACAACAGAAAAATATGTAATTTAAAAATTTGTTGAA	5580
Qy	870	AGGAGGTACCTCTCATTTCTTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCAA	929
Db	5581	AGGAGGTACCTCTCATTTCTTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCAA	5640
Qy	930	TATTTTATATATGTAAGTTTATTTTATTAAGTATACATTTTATTTATGTGAGTTTATTA	989
Db	5641	TACTTTATATATGTAAGTTTATTTTATTAAGTATACATTTTATTTATGTGAGTTTATTA	5700
Qy	990	ATATGGAATTTTATTAAGAAACATTTCTGCTATTGATATTTT-AGTATAAGGCAAAATAT	1048
Db	5701	ATATGGAATTTTATTAAGAAACATTTCTGCTATTGATATTTTGTAGTATAAAGCAAAATAT	5760
Qy	1049	ATTATGACATAACTATGGAACAAGATATCTTAGGCTTTTAAATAACACATGTGATATCA	1108
Db	5761	ATTATGATAATAACTATGGAACAAGATATCTTAGGCTTTTAAATAACACATGATATCA	5820
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Db	5821	TAAA 5824	
RESULT 5			
US-11-102-240-153			
; Sequence 153, Application US/11102240			
; Publication No. US20050260647A1			
; GENERAL INFORMATION:			
; APPLICANT: Goddard, Audrey			
; APPLICANT: Godowski, Paul			
; APPLICANT: Grimaldi, Christopher J.			
; APPLICANT: Gurney, Austin L.			
; APPLICANT: Wood, William I.			
; TITLE OF INVENTION: ANTIBODIES TO POLYPEPTIDES ENCODED BY A NUCLEIC ACID UNDEREXPRESS			
; TITLE OF INVENTION: ESOPHAGEAL TUMOR			
; FILE REFERENCE: P3230R1C106C			
; CURRENT APPLICATION NUMBER: US/11/102,240			
; CURRENT FILING DATE: 2005-04-08			
; PRIOR APPLICATION NUMBER: 10/063662			
; PRIOR FILING DATE: 2002-05-07			
; PRIOR APPLICATION NUMBER: 10/006867			
; PRIOR FILING DATE: 2001-12-06			
; PRIOR APPLICATION NUMBER: PCT/US00/23328			
; PRIOR FILING DATE: 2000-08-24			
; PRIOR APPLICATION NUMBER: 60/170262			
; PRIOR FILING DATE: 1993-12-09			
; NUMBER OF SEQ ID NOS: 170			
; SEQ ID NO 153			
; LENGTH: 1152			
; TYPE: DNA			
; ORGANISM: Homo Sapien			
US-11-102-240-153			
Query Match 46.9%; Score 524.8; DB 7; Length 1152;			
Best Local Similarity 73.2%; Pred. No. 3e-110;			
Matches 832; Conservative 0; Mismatches 267; Indels 37; Gaps 11;			
Qy	9	CTCTCCTCTCACTTATCAACTGTTGACACTGTGTGGGATCTCTGATGGCTCTCTGCGAGAA	68
Db	15	CTCCTTCCCAGTCACCAAGTTGCTCGAGTTAGAAATGTCTGCAATGGCGCCCTGCGAGAA	74
Qy	69	ATCTATGAGTTTTCCTCTATGGGACTTTTGGCCGCCAGCTTGCCTGCTTCTCATTTGCCCT	128

RESULT 6

Db	75	ATCTGTGAGCTCTTTTCTTATGGGACCTTGGCCACCAGCTGCTCTTCTCTTGTGGCCCT	134
Qy	129	GTGGGCCAGGAGGAAATGCGCTGCCCTCAACACCCGGTCAAGCTTGGGTGTCCAA	188
Db	135	CTTGGTACAGGGAGGAGCTGCGCCCATCAGCTCCCACTGAGGGTTTGACAACTCAA	194
Qy	189	CTTCCAGCAGCGCTACATCGTCAACCGCACCTTTATGCTGGCCCAAGGAGGCGAGCTTGC	248
Db	195	CTTCCAGCAGCGCTTATATACCAACCGCACCTTATGCTGGCTTAAGGAGGCTAGCTTGC	254
Qy	249	AGATAACAACACAGACGCTCCGGCTCATCGGGGAGAACTGTTCCGAGGAGTCAAGTGTAA	308
Db	255	TGATAACAACACAGACGCTTCTCTCATTTGGGGAGAACTGTTCCACGAGTCAAGTATGAG	314
Qy	309	AGATCAGTGTCTACTGATGAGCAGGTGCTCACTTCACTCCCTGGAGAGCTTCTGCTCCC	368
Db	315	TGAGCGCTGCTATCTGATGAAGCAGGTGCTGAACTTCACTTGAAGAGTGTGTTCCC	374
Qy	369	CCAGTCAGACAGGTTCCAGCCCTACATGAGGAGGTGGTACCTTTCTGACCAAACTCAG	428
Db	375	TCAATCTGATAGGTTCCAGCTTATATGAGGAGGTGGTGGCTTCTTGGCCAGGCTCAG	434
Qy	429	CAATCAGCTCAGCTCCTGTCACTCAGCGGTGACGACCAACAATCCAGAGAAATGTGCA	488
Db	435	CAACAGGCTAAGCACATGTCTATTTGAAGGTGATGACCTGCATATCCAGAGGAATGTGCA	494
Qy	489	AGGCTGAAGGAGACAGTGAAGAAAGCTTGGAGAGGTGAGAGATCAAGCGATTTGGGA	548
Db	495	AAAGCTGAAGGACACAGTGAAGAAAGCTTGGAGAGGTGAGAGATCAAGCAATTTGGAGA	554
Qy	549	ACTGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAACTAGAA	608
Db	555	ACTGGATTTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAACTAGAA	614
Qy	609	AACGAAGAACTGCTTCTTCTGCTTCTTAAAGAAACAATAAGATCCCTGAATGGACTTT	668
Db	615	AATGAATAACTAACCCCTTTTCCCTGCTAGAAATAACAATTAGATGCTCCCAAGCGATTT	674
Qy	669	TTTA-----CTAAGGAAGTGAGAGCTAAGCTCCATCATCTAGAGATTTTCAATGA	724
Db	675	TTTTTAAACCAAGGAAGATGGGGAAGCAACTCCATCATGAGGTGGTGGATTTCAAATGA	734
Qy	725	AACCTGCTCAGTTTGAAGAAAGAAATAGTGTCAA--GTTGTCCATGAGACGAG--AGGTAG	781
Db	735	ACCCTGCTGTTAGTTACAAAGGAACCAATGCCACTTTTGTTTATAGACCAAGAGGTAG	794
Qy	782	ACTTGATAACCAACAAAGATTCATTGACAAATATTTTATGTCACTGATG---ATACAACA	837
Db	795	ACTTTCTAAGCATAGATATTTATTTGATAACATTTTCATTGTAACCTGTTCTATACACAG	854
Qy	838	GAAAAATAATGTACTTTTAAAAAATTTGTTT-----GAAAGGAGTTTACCTCTCATTCCT	890
Db	855	AAAACAATTTATTTTTTAAAAATAATTTGCTTTTCCATATAAAAAAGATTACTTTCATTCCT	914
Qy	891	TTA---GAAAAAGAGCTTATGTAACTTCA--TTTCCATATCCAATATTTTATATATGTAA	945
Db	915	TTAGGGGAAAAAACCCTTAATAGCTTCATGTTTCCATATCATCTACTTTATATTTATAA	974
Qy	946	GTTTATTTTATTAAGTATA-----CATTTTATTTATGTCAGTTTATTAATATGGAATTT	999
Db	975	ATGTATTTTATTTATTAAGACTGCAATTTTATTTATATCATTTTATTAATATGGAATTT	1034
Qy	1000	ATTTATAGAAACATATCTGCTATTGATA--TTTAGTATAGGCAATAA---ATATTTATG	1055
Db	1035	ATTTATAGAAACATCATTTCCGATATTTGCTAGTGTAGTGTAGGCTAAATTTATTTATG	1094
Qy	1056	ACAATAACTATGG---AAACAAGATATCTTAGGCTTTAATAAACAACATGATGATATC	1107
Db	1095	ACAATAATTTATAGAGCTATTAACATGTTTATTTTGACCTCAATAAACACTTGGATATC	1150

Db 629 AATGAATAACTAACCCCTTTCCCTCTAGAAATAACAATTAGATGCCCAAAGCGATTT 688

Qy 669 TT 670  
||

Db 689 TT 690

RESULT 7  
US-11-177-987-18  
; Sequence 18, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Indu  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 18  
; LENGTH: 418  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-11-177-987-18

Query Match 20.0%; Score 223.6; DB 7; Length 418;  
Best Local Similarity 74.2%; Pred. No. 6.9e-42;  
Matches 297; Conservative 0; Mismatches 99; Indels 4; Gaps 1;

Qy 354 AGACGTTCTGCTCCCCAGTCCAGACAGGTTCCAGGCCTTACATGCAGGAGTGGTACCTTT 413  
Db 1 AGAAGTGCTGTTCCCTCAATCTGATAGGTTCCAGCCTTATATGCAGGAGTGGTGCCCTT 60

Qy 414 CCTGACCAAACTCAGCAATCAGCTCAGCTCCTGTCAATCATCAGCGGTGACGACAGAACAT 473  
Db 61 CCTGGCCAGCTCAGCAACAGGCCTAAGCACATGTCTATATTGAAGGTGATGACCTGCATAT 120

Qy 474 CCAAGAATGTGAGAAGGCTGAAGAGACAGTGAAGAAAGCTTGGAGAGAGTGGAGAGAT 533  
Db 121 CCAGAGGAATGTGCAAAAGCTGAAGGACACAGTGAAGAAAGCTTGGAGAGAGTGGAGAGAT 180

Qy 534 CRAAGCGATTGGGGAACTGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGC 593  
Db 181 CAAGCAATTTGGAAGAACTGGATTGCTGTTATGCTCTGAGAAATGCTCTGAGAAATGCTCTGAGC 240

Qy 594 GAGAAGAAGCTGAGAAACGAAGAACTGCTCCTTCTGCTCCTTTCTTAAAGAAACAAATAAGAT 653  
Db 241 AGAGCAAGCTGAAATAATGAATACTAACCCCTTTCCCTGCTAGAAATAACAATTAGAT 300

Qy 654 CCTGTAATGACTTTTTT-----ACTAAGGAAAGTGAAGGCTTAACGTCCATCATCATTA 709  
Db 301 GCCCCAAAGCGATTTTTTTTTTAAACCAAAAGGAAGATGGGAAGCCAAACTCCATCATGATGG 360

Qy 710 GAAGATTTCACATGAACCTGGCTCAGTTCAGAAAGAAA 749  
Db 361 GTGGATTCCAAATGAACCCCTGCGTTAGTTACAAAGGAAA 400

RESULT 8  
US-11-177-987-26  
; Sequence 26, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:

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; APPLICANT: Dumoutier, Laure
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 26
; LENGTH: 4797
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-11-177-987-26

Query Match      11.3%; Score 126; DB 7; Length 4797;
Best Local Similarity 71.7%; Pred. No. 3.6e-19;
Matches 165; Conservative 0; Mismatches 65; Indels 0; Gaps 0;

Qy      9 CTCTCCTCTCACTTATCAACTGTGACACTGTGCGGATCTCTGATGCTGCTGCTGCGAGAA 68
Db      29 CTCCTTCCCCAGTCACCACTGTGCGGATTTGCTGAGATTGCTGCAATGGCCGCCCTGCGAGAA 88

Qy      69 ATCTATGAGTTTTCCCTTATGGGACCTTTGGCCGCCAGCTGCTGCTGCTCTCTCATTTGCCCT 128
Db      89 ATCTGTGAGCTTCTTCTTATGGGACCCCTGGCCACACAGCTGCTCTCTCTTGGCCCT 148

Qy      129 GTGGGCCAGAGGAGCAATGCGCTGCGCGTCAACACCCGGTGCAGCTTGAGGTGTCCAA 189
Db      149 CTTGGTACAGGAGGAGCAGCTGCGGCCCATCAGCTCCCACTGCGAGGCTTGACAAGTCCAA 208

Qy      189 CTTCCAGACGCGTACATCGTCAACCGCACCTTTATCTGCTGCGCAAGGAGG 238
Db      209 CTTCCAGACGCGCTATATACCAACCGCACCTTTATCTGCTGCGCAAGGAGG 258

RESULT 9
US-10-995-561-13313/c
; Sequence 13313, Application US/10995561
; Publication No. US20050272054A1
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
; TITLE OF INVENTION: DETECTION AND USES THEREOF
; FILE REFERENCE: CL001559
; CURRENT APPLICATION NUMBER: US/10/995,561
; CURRENT FILING DATE: 2004-11-24
; NUMBER OF SEQ ID NOS: 85702
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13313
; LENGTH: 26772
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-995-561-13313

Query Match      4.9%; Score 54.6; DB 6; Length 26772;
Best Local Similarity 52.4%; Pred. No. 0.014;
Matches 140; Conservative 1; Mismatches 125; Indels 1; Gaps 1;

Qy      852 TTTAAAAAATGTTGAAGAGGGTTACCTCTCATCTCTTTAGAAAAAAGCTTATCTGTA 911
Db      21973 TCTCAAAATACATATATAATAATTTATATAATAAATTTCTATATAAAATTTATATAT 21914

Qy      912 CTTCAATTTCCCATATCCCAATATTTTATATATGTAAGTTTATTTATATAAGTATACATTTT 971

US-10-995-561-13479/c
; Sequence 13479, Application US/10995561
; Publication No. US20050272054A1
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
; TITLE OF INVENTION: DETECTION AND USES THEREOF
; FILE REFERENCE: CL001559
; CURRENT APPLICATION NUMBER: US/10/995,561
; CURRENT FILING DATE: 2004-11-24
; NUMBER OF SEQ ID NOS: 85702
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13479
; LENGTH: 54946
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-995-561-13479

Query Match      4.9%; Score 54.6; DB 6; Length 54946;
Best Local Similarity 52.4%; Pred. No. 0.02;
Matches 140; Conservative 1; Mismatches 125; Indels 1; Gaps 1;

Qy      852 TTTAAAAAATGTTGAAGAGGGTTACCTCTCATCTCTTTAGAAAAAAGCTTATCTGTA 911
Db      3427 TCTCAAAATACATATATAATAATTTATATAATAAATTTCTATATAAAATTTATATATAT 3368

Qy      912 CTTCAATTTCCCATATCCCAATATTTTATATATGTAAGTTTATTTATATAAGTATACATTTT 971
Db      3367 ATTATATATAATTTCTATATAAAATATATATATATATATAATAAT-ATTCTATATAAA 3309

Qy      972 ATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAACATTTCTGCTATTCGATTTT 1031
Db      3308 ATATATGTTATTTATTTATAGAAATCTATATAATAATTTATATAATAATTTATATATATTA 3249

Qy      1032 AGTATAGGCAATAATATTTATGACAATACTATGGAACAAGATATCTTAGCGCTTTAA 1091
Db      3248 TATATTATAATTAATAATTTATATAATGATTTATATAATAATTTATATAATAATTAATAATA 3189

Qy      1092 TAAACACATGGAATCATCAAAAAAAA 1118
Db      3188 TTTATATTATATAATTTATATAATAA 3162

RESULT 11
US-11-121-086-105
; Sequence 105, Application US/11121086
; Publication No. US20050266459A1
; GENERAL INFORMATION:
; APPLICANT: POULSEN, TIM S.
; APPLICANT: NIELSEN, KIRSTEN V.
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES
; FILE REFERENCE: 09138.6000-00000
; CURRENT APPLICATION NUMBER: US/11/121,086
; CURRENT FILING DATE: 2005-05-04
; PRIOR APPLICATION NUMBER: 60/567,570

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Db      21913 ATTATATAATAATTTCTATATAAAATATATATATTTATATAATAAT-ATTCTATATAAA 21855
Qy      972 ATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAACATTTCTGCTATTCGATTTT 1031
Db      21854 ATATATGTTATATTTATTTATAGAAATCTATATAATAATTTATATAATAATTTATATATTA 21795
Qy      1032 AGTATAGGCAATAATAATTTATGACAATACTATGGAACAAGATATCTTAGCGCTTTAA 1091
Db      21794 TATATTATAATTAATAATTTATATAATGATTTTATATAATAATTTATATAATAATAATA 21735
Qy      1092 TAAACACATGGAATCATCAAAAAAAA 1118
Db      21734 TTTATATTATATAATTTATATAATAA 21708

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RESULT 10
US-10-995-561-13479/c
; Sequence 13479, Application US/10995561
; Publication No. US20050272054A1
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
; TITLE OF INVENTION: DETECTION AND USES THEREOF
; FILE REFERENCE: CL001559
; CURRENT APPLICATION NUMBER: US/10/995,561
; CURRENT FILING DATE: 2004-11-24
; NUMBER OF SEQ ID NOS: 85702
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13479
; LENGTH: 54946
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-995-561-13479

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Query Match      4.9%; Score 54.6; DB 6; Length 54946;
Best Local Similarity 52.4%; Pred. No. 0.02;
Matches 140; Conservative 1; Mismatches 125; Indels 1; Gaps 1;

Qy      852 TTTAAAAAATGTTGAAGAGGGTTACCTCTCATCTCTTTAGAAAAAAGCTTATCTGTA 911
Db      3427 TCTCAAAATACATATATAATAATTTATATAATAAATTTCTATATAAAATTTATATATAT 3368

Qy      912 CTTCAATTTCCCATATCCCAATATTTTATATATGTAAGTTTATTTATATAAGTATACATTTT 971
Db      3367 ATTATATATAATTTCTATATAAAATATATATATATATATAATAAT-ATTCTATATAAA 3309

Qy      972 ATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAACATTTCTGCTATTCGATTTT 1031
Db      3308 ATATATGTTATTTATTTATAGAAATCTATATAATAATTTATATAATAATTTATATATATTA 3249

Qy      1032 AGTATAGGCAATAATATTTATGACAATACTATGGAACAAGATATCTTAGCGCTTTAA 1091
Db      3248 TATATTATAATTAATAATTTATATAATGATTTATATAATAATTTATATAATAATTAATAATA 3189

Qy      1092 TAAACACATGGAATCATCAAAAAAAA 1118
Db      3188 TTTATATTATATAATTTATATAATAA 3162

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RESULT 11
US-11-121-086-105
; Sequence 105, Application US/11121086
; Publication No. US20050266459A1
; GENERAL INFORMATION:
; APPLICANT: POULSEN, TIM S.
; APPLICANT: NIELSEN, KIRSTEN V.
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES
; FILE REFERENCE: 09138.6000-00000
; CURRENT APPLICATION NUMBER: US/11/121,086
; CURRENT FILING DATE: 2005-05-04
; PRIOR APPLICATION NUMBER: 60/567,570

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APPLICANT: CARGILL, Michele et al.  
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH  
CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF  
DETECTION AND USES THEREOF  
FILE REFERENCE: C1001559  
CURRENT APPLICATION NUMBER: US/10/995,561  
CURRENT FILING DATE: 2004-11-24  
NUMBER OF SEQ ID NOS: 85702  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 13354  
LENGTH: 47444  
TYPE: DNA  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: misc.feature  
LOCATION: (1)...(47444)  
OTHER INFORMATION: n = A,T,C or G, or insertion/deletion polymorphism (see Tables 1-  
US-10-995-561-13354

Query Match 4.8%; Score 53.4; DB 6; Length 47444;  
Best Local Similarity 49.8%; Pred. No. 0.036;  
Matches 163; Conservative 0; Mismatches 161; Indels 3; Gaps 1;  
Qy 793 ACAAGATTTCATGACAAATATTTTATGTCATGATGATACACAGAAAAAATATGTAAT 852  
Db ATATATAATTATGATGATACATAATATACATAATATATATATATATATATATATATATAT 27574  
Qy 853 TTAAGAAATGTTGAAGAGGTTACCTCTCATCTCCCTTTAGAAAAAAGCTTATGTAAC 912  
Db ATATACATATAATATAAT 27634  
Qy 913 TTCAATTCATATCCAAATATTTTATATATATGTAAGTTTATTTTATTAAGTATACATTTTA 972  
Db TATGATACAT 27694  
Qy 973 TTTATGTCAGTTTATTAATATGAGTTTATTTATAGAAACATTTCTGCTATGATATTTTA 1032  
Db TATATAATATATATGATATACATAATATACATAATATATATATATATATATATATATATAT 27754  
Qy 1033 GTATAAGGCAAAATATATTTATGACATAATCTATGGAACAAAGATATCTTAGGCTTTAAT 1092  
Db ATATA---CATATAATATATATATATATGATACATAATATATATATATATATATATATATA 27811  
Qy 1093 AACACATGGATATCATAAABAAAA 1119  
Db TATTATGATACATAATATATATATAT 27838

RESULT 15  
US-11-121-086-25/c  
Sequence 25, Application US/11121086  
Publication No. US20050266459A1  
GENERAL INFORMATION:  
APPLICANT: NIELSEN, KIRSTEN V.  
TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES  
FILE REFERENCE: 09138.6000-00000  
CURRENT APPLICATION NUMBER: US/11/121,086  
CURRENT FILING DATE: 2005-05-04  
PRIOR APPLICATION NUMBER: 60/567,570  
PRIOR FILING DATE: 2004-05-04  
NUMBER OF SEQ ID NOS: 107  
SOFTWARE: PatentIn version 3.3  
SEQ ID NO 25  
LENGTH: 173602  
TYPE: DNA  
ORGANISM: Homo sapiens  
US-11-121-086-25

Query Match 4.7%; Score 52.8; DB 7; Length 173602;  
Best Local Similarity 51.7%; Pred. No. 0.089;  
Matches 153; Conservative 0; Mismatches 132; Indels 11; Gaps 1;

Qy 799 ATTCAATTGCAATATATTTTATTTGTCACATGATGATACACAGAAAAAATATGTAATTTAAAA 858  
Db ATATATTTATATATAATTTTATATATATTTGATATATATTTATATATATATATATATATATAT 136327  
Qy 859 AATTGTTTGAAGAGGTTTACCTCTCATCTCCCTTTAGAAAAAAGCTTATGTAACCTTCATT 918  
Db ATAAATATATAAATATATATAATTTTATATATATATATATATATATATATATATATATATAATT 136267  
Qy 919 TCCATATCCAATATTTTATATATATGTAAGTTTATTTTATTAAGTATATACATTTTATTTATG 978  
Db TATATAATTTTA 136207  
Qy 979 TCAGTTTATTAATATGGAATTTATTTATAGAAACATTTCTGCTATGATATTTAGTATAA 1038  
Db T-----TATATATTTAA 136158  
Qy 1039 GGCATAATATATTTATGACAAATTAATCTATGGAACAAAGATATCTTAGGCTTTAATAA 1094  
Db TATATATTTATATATATATATATATTTTATAAATATATATATATATATATATATATATATAA 136102

Search completed: December 21, 2005, 18:11:02  
Job time : 82.0229 secs



GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 842.71 Seconds  
(without alignments)  
15704.028 Million cell updates/sec

Title: US-09-751-797-8

Perfect score: 7445

Sequence: 1 gctatcacctgcttaagat.....gattaattaattatgtgat 7445

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 1303057 seqs, 888780828 residues

Total number of hits satisfying chosen parameters: 2606114

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued Patents NA.\*

- 1: /cgn2\_6/ptodata/1/ina/1\_COMB.seq.\*
- 2: /cgn2\_6/ptodata/1/ina/5\_COMB.seq.\*
- 3: /cgn2\_6/ptodata/1/ina/6A\_COMB.seq.\*
- 4: /cgn2\_6/ptodata/1/ina/6B\_COMB.seq.\*
- 5: /cgn2\_6/ptodata/1/ina/H\_COMB.seq.\*
- 6: /cgn2\_6/ptodata/1/ina/PTUS\_COMB.seq.\*
- 7: /cgn2\_6/ptodata/1/ina/PP\_COMB.seq.\*
- 8: /cgn2\_6/ptodata/1/ina/RE\_COMB.seq.\*
- 9: /cgn2\_6/ptodata/1/ina/backfiles1.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	7445	100.0	7445	3	US-09-178-973B-8
2	7445	100.0	7445	3	US-09-419-568F-8
3	7445	100.0	7445	3	US-09-354-243B-8
4	4245.2	57.0	5935	3	US-09-178-973B-17
5	4245.2	57.0	5935	3	US-09-419-568F-29
6	4245.2	57.0	5935	3	US-09-354-243B-29
7	1081	14.5	8888	3	US-09-949-016-17185
8	686	9.2	4797	3	US-09-419-568F-25
9	686	9.2	4797	3	US-09-354-243B-25
10	601.4	8.1	1119	3	US-09-178-973B-7
11	601.4	8.1	1119	3	US-09-419-568F-7
12	601.4	8.1	1119	3	US-09-354-243B-7
13	598.2	8.0	1166	3	US-10-084-298-3
14	555.2	7.5	1111	3	US-09-178-973B-9
15	555.2	7.5	1111	3	US-09-419-568F-9
16	555.2	7.5	1111	3	US-09-354-243B-9
17	499	6.7	1050	3	US-10-090-365-40
18	217.6	2.9	1191	3	US-10-084-298-1
19	215.6	2.9	1116	3	US-10-090-365-14
20	215.6	2.9	1116	3	US-09-728-911-14
21	214.2	2.9	1152	3	US-09-870-574-1
22	181.4	2.4	601	3	US-09-949-016-190092
23	126	1.7	689	3	US-09-949-016-5443
24	126	1.7	690	3	US-09-419-568F-24

## ALIGNMENTS

## RESULT 1

US-09-178-973B-8

; Sequence 8, Application US/09178973B

; Patent No. 6274710

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Louhed, Jamila

; APPLICANT: Renauld, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; TITLE OF INVENTION: (TIFFS)

; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5543

; CURRENT APPLICATION NUMBER: US/09/178,973B

; CURRENT FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 17

; SEQ ID NO 8

; LENGTH: 7445

; TYPE: DNA

; ORGANISM: Mus musculus

US-09-178-973B-8

Query Match 100.0%; Score 7445; DB 3; Length 7445;

Best Local Similarity 100.0%; Pred. No. 0;

Matches 7445; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	GTCTATCACTGCTTAAGATTCTTCTAAATTTATAAAAAAACTATTCTTAAATGAAAA	60
Db	1	GTCTATCACTGCTTAAGATTCTTCTAAATTTATAAAAAAACTATTCTTAAATGAAAA	60
Qy	61	GCAACAGAGCAGTATTTATAGCATGGTCTTGACCATGCGAGGTACAGAGTGGATGG	120
Db	61	GCAACAGAGCAGTATTTATAGCATGGTCTTGACCATGCGAGGTACAGAGTGGATGG	120
Qy	121	TAAGAGCGCTATATCAGCATTTAAACAATGTTTAAATGTTTCTTCTGCGAAGCAAACT	180
Db	121	TAAGAGCGCTATATCAGCATTTAAACAATGTTTAAATGTTTCTTCTGCGAAGCAAACT	180
Qy	181	TGAATCTATGCTTAAACAATCTTCAAGCCTCTATATAGTCTACGACTGAGTCCG	240
Db	181	TGAATCTATGCTTAAACAATCTTCAAGCCTCTATATAGTCTACGACTGAGTCCG	240
Qy	241	CTGCTGTCAACAGAGCTCTTGAGCAGCTCTCTCTGTTTGAATTTATGTTCTTTGA	300
Db	241	CTGCTGTCAACAGAGCTCTTGAGCAGCTCTCTCTGTTTGAATTTATGTTCTTTGA	300
Qy	301	TCGACTCCCAACCTCTCACTTTCGGTCTCTGATGGCCACTTTCACCTTTCGCAATTA	360





Db 4681 AAAGTATGAGATGGAAACCCCTTCTTACCTTTTACCTTCATTTCTTAGTTTTTTTTTTC 4740  
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Db  
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Db  
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Db  
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Db  
6421 AGAAGGCTGATFACCTGGTTTTGGTGTCTCAGCAAGCAAAATGTCACAGCTCTTTCTAACTG 6480  
Qy  
6481 GTACCACTTTAGAAAAATGCTGCTCAAAATGGTTGATTAATCTTTTTCATAGCT 6540  
Db  
6481 GTACCACTTTAGAAAAATGCTGCTCAAAATGGTTGATTAATCTTTTTCATAGCT 6540  
Qy  
6541 TGGAGAGAGTGGAGAGATCAAGGCGATTGGGAACTGGACCTGCTGTTTATGTCTCTGAG 6600  
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6781 CAAGTTGTCCATGAGACCAGAGGTAGACTTGTATAACCAAAAGATTCATTCAATATTT 6840  
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Db  
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Qy	1141	TTGCAAGAAGTAAATGTCTAGAGAAATAGCAAAATGTATAGTATTATTTATTTTAAAAAAA	1200
Db	1141	TTGCACAAGTAAAAATGTCTAGAGAAATAGCAAAATGTATAGTATTATTTATTTTAAAAAAA	1200
Qy	1201	TCTATGCTTTAAAATGCTATTATAGATTGCTTCACATACCGATATTTCCAAACCTTAACCTTGACC	1260
Db	1201	TCTATGCTTTAAAATGCTATTATAGATTGCTTCACATACCGATATTTCCAAACCTTAACCTTGACC	1260
Qy	1261	TTGGCTATGATTTTCAACCTTTGTATTTTGGCATCTACCATAAACAGTCTCTGGAACAGAAACAT	1320
Db	1261	TTGGCTATGATTTTCAACCTTTGTATTTTGGCATCTACCATAAACAGTCTCTGGAACAGAAACAT	1320
Qy	1321	TCTGTGGCAATGGGAGCTGTGAAGAAAGCCAAATCTTATTAATAAAAAAAGAAACAGCTA	1380
Db	1321	TCTGTGGCAATGGGAGCTGTGAAGAAAGCCAAATCTTATTAATAAAAAAAGAAACAGCTA	1380
Qy	1381	GTTATAGTTTAGGATTCATATACTATAAAAAATAGAGATATAATATTTTAAATTTGA	1440
Db	1381	GTTATAGTTTAGGATTCATATACTATAAAAAATAGAGATATAATATTTTAAATTTGA	1440
Qy	1441	AATAATCTCCAAGTTTTCATTATGGCTTATTTCAAAGCACAGAATATAGGACACGGGTCT	1500
Db	1441	AATAATCTCCAAGTTTTCATTATGGCTTATTTCAAAGCACAGAATATAGGACACGGGTCT	1500
Qy	1501	TTTATTTCTGTGTCATTTCTAAAGAGATAAGAATCTATGAATGGTGGGAAAATAGTCTC	1560
Db	1501	TTTATTTCTGTGTCATTTCTAAAGAGATAAGAATCTATGAATGGTGGGAAAATAGTCTC	1560
Qy	1561	GTGACCAAAACGCTGACTCAATAGCTACGGGAGATCAAGGCTGCTCTACTCAATCAGAA	1620
Db	1561	GTGACCAAAACGCTGACTCAATAGCTACGGGAGATCAAGGCTGCTCTACTCAATCAGAA	1620
Qy	1621	TCTACTACGGCAAAAGCCATGGCTTCTTTGAAAAACCGTGTTTAGAAGATTTCTGGGATTT	1680
Db	1621	TCTACTACGGCAAAAGCCATGGCTTCTTTGAAAAACCGTGTTTAGAAGATTTCTGGGATTT	1680
Qy	1681	GTGTGCAAAAGCACCTTTGTGGCCCTCACCGTGACGTTTTTAGGGAAGACTTCCCATCTCT	1740
Db	1681	GTGTGCAAAAGCACCTTTGTGGCCCTCACCGTGACGTTTTTAGGGAAGACTTCCCATCTCT	1740
Qy	1741	CAAGGTGGGAAGGCTTGGAGGTGTGTCTGTGTGGCCCTCCTATGTTGGTTAGGTACTTCTC	1800
Db	1741	CAAGGTGGGAAGGCTTGGAGGTGTGTCTGTGTGGCCCTCCTATGTTGGTTAGGTACTTCTC	1800
Qy	1801	AGAGACAGGACTGGAATTAGATAATGTCTGATGTCTATCATTCATCAATACCAAAAAA	1860
Db	1801	AGAGACAGGACTGGAATTAGATAATGTCTGATGTCTATCATTCATCAATACCAAAAAA	1860
Qy	1861	ACCTGTGTCCGATGGCTATAAAGCAGCAACTTCTGCCTCTCCCATCACAGCAGAG	1920
Db	1861	ACCTGTGTCCGATGGCTATAAAGCAGCAACTTCTGCCTCTCCCATCACAGCAGAG	1920
Qy	1921	ACACCTAAACAGGTAAGCACTCAGACCTCTACAGACAATCATCTGCTTGGTACCATGCTA	1980
Db	1921	ACACCTAAACAGGTAAGCACTCAGACCTCTACAGACAATCATCTGCTTGGTACCATGCTA	1980
Qy	1981	CCGACGAACTGCTCCCTCATGTTTTTGCTTTTGCCTCTCTCACTAAACAGGCTCTCCT	2040
Db	1981	CCGACGAACTGCTCCCTCATGTTTTTGCTTTTGCCTCTCTCACTAAACAGGCTCTCCT	2040
Qy	2041	CTCACTTATCAACTGTTGACACTTGTGCGAATCTCTGATGGCTGCTCCGACAGAAATCTATG	2100
Db	2041	CTCACTTATCAACTGTTGACACTTGTGCGAATCTCTGATGGCTGCTCCGACAGAAATCTATG	2100
Qy	2101	AGTTTTTCCCTTATGGGACCTTTGGCCGCCAGTGCCTGTCTTCTCATTTGCCCTGTGGGCC	2160
Db	2101	AGTTTTTCCCTTATGGGACCTTTGGCCGCCAGTGCCTGTCTTCTCATTTGCCCTGTGGGCC	2160
Qy	2161	CAGGAGCAATGCGCTGCCCGTCAACACCCGGTGCAAGCTTGAGGTGTCCAATCTCCAG	2220
Db	2161	CAGGAGCAATGCGCTGCCCGTCAACACCCGGTGCAAGCTTGAGGTGTCCAATCTCCAG	2220
Qy	2221	CAGCCGTATCATCTGTCAACCGCACCTTTATGTCTGGCCAGGAGGTACAGCTGCATCTCTTT	2280

Db	2221	 CAGCGTACATGTCAAACGACCTTTATGCTGGCCAGAGGTA	2280	 CAGCTGATCTCTTT
Qy	2281	CTCTCCATACCGCTTTGCCATTTTCTCTGAAGACACTTGC	2340	CTTTTAGGGCGCTTTA
Db	2281	CTCTCCATACCGCTTTGCCATTTTCTCTGAAGACACTTGC	2340	CTTTTAGGGCGCTTTA
Qy	2341	CTCTCGGAGGTCACATACCTATGTTTTCTCTCTTTAGAGAC	2400	CTTTTAAGGACTGGGT
Db	2341	CTCTCGGAGGTCACATACCTATGTTTTCTCTCTTTAGAGAC	2400	CTTTTAAGGACTGGGT
Qy	2401	CTTTTCTATTTCTATTTCAAGGCTCTCAGACCATTTCTAT	2460	CTTTCTGGCTTACGACACA
Db	2401	CTTTTCTATTTCTATTTCAAGGCTCTCAGACCATTTCTAT	2460	CTTTCTGGCTTACGACACA
Qy	2461	TATACTGAATTTTATCTACAGAGCGCATTTAGAAAGCCAC	2520	CTTTGCAATATCTTTC
Db	2461	TATACTGAATTTTATCTACAGAGCGCATTTAGAAAGCCAC	2520	CTTTGCAATATCTTTC
Qy	2521	CAATTCCTGTCTCTCTCTGAACTCATCTCTCTTGGCTAC	2580	CTCTGAGACCACTGGG
Db	2521	CAATTCCTGTCTCTCTCTGAACTCATCTCTCTTGGCTAC	2580	CTCTGAGACCACTGGG
Qy	2581	GACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTT	2640	CTTTTCTGCTTCTGCT
Db	2581	GACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTT	2640	CTTTTCTGCTTCTGCT
Qy	2641	TTTCTCTCTTTTCAGGCGAGCTTTCAGAGTAACAAACAC	2700	CTTCGGGGAGA
Db	2641	TTTCTCTCTTTTCAGGCGAGCTTTCAGAGTAACAAACAC	2700	CTTCGGGGAGA
Qy	2701	AACCTGTTCCGAGGAGTCAGTGAAGTCTCTACCTGTGAT	2760	CTGAGGAGGCTAGCTCGGGAGC
Db	2701	AACCTGTTCCGAGGAGTCAGTGAAGTCTCTACCTGTGAT	2760	CTGAGGAGGCTAGCTCGGGAGC
Qy	2761	TGTTGGACCTCTGGGATAGTCTGACGTATGACCCCTGCT	2820	CTTGTCTACTCCTCAGG
Db	2761	TGTTGGACCTCTGGGATAGTCTGACGTATGACCCCTGCT	2820	CTTGTCTACTCCTCAGG
Qy	2821	CTAAAGATCAGTGTCTACTCTGATGAAGCAGAGTGCTCA	2880	CTTCACTTCAAGAGAGGCTTCTGC
Db	2821	CTAAAGATCAGTGTCTACTCTGATGAAGCAGAGTGCTCA	2880	CTTCACTTCAAGAGAGGCTTCTGC
Qy	2881	TCCTCCAGTCAGACAGGTTCCAGCCCTACATGAGAGAGTG	2940	CTTTCCTGACCAAAAC
Db	2881	TCCTCCAGTCAGACAGGTTCCAGCCCTACATGAGAGAGTG	2940	CTTTCCTGACCAAAAC
Qy	2941	TCAGCAATCAGCTCAGCTCCTCTGTGTAAGTCTGACTCT	3000	CTGATGCTCCTCTCTCTT
Db	2941	TCAGCAATCAGCTCAGCTCCTCTGTGTAAGTCTGACTCT	3000	CTGATGCTCCTCTCTCTT
Qy	3001	CCTCTTCTATTTCAGATGAAGAACCGAGGTCCTGCCCC	3060	CTTCTTCAAGAGTGAAGGA
Db	3001	CCTCTTCTATTTCAGATGAAGAACCGAGGTCCTGCCCC	3060	CTTCTTCAAGAGTGAAGGA
Qy	3061	GGGCTTCAGCACCAACCATCATAGGCCACTTGAATAGGT	3120	CTCAAGGCTTTGGCTTC
Db	3061	GGGCTTCAGCACCAACCATCATAGGCCACTTGAATAGGT	3120	CTCAAGGCTTTGGCTTC
Qy	3121	AATTGAGTAATACTTTGAGTTTGTATGAGTGAAGCTTTA	3180	TTTGTATTCATCTGGAAGA
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Qy	3181	AATCAACTCAAAATCTCTGAGATGAGAAAGATGTTGGG	3240	CTAAAGGCTTAGATAGA
Db	3181	AATCAACTCAAAATCTCTGAGATGAGAAAGATGTTGGG	3240	CTAAAGGCTTAGATAGA
Qy	3241	GAAGACAGATCTGCTGAGTATAGTACTTTATGGGGGAGC	3300	CTAGGCGGATATCCACTGAGTA
Db	3241	GAAGACAGATCTGCTGAGTATAGTACTTTATGGGGGAGC	3300	CTAGGCGGATATCCACTGAGTA
Qy	3301	CAAGTACTTGTGGGAGAGAAATCCACTGAGTACAGTACT	3360	TTGTTGCACTGGAGATCCAC



Db 3301 CAAGTACTTGTGGGGAGAGAAATCCACTGTAGTCAAGTACTTGTGTGCATGGAGATCCAC 3360  
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Db 3361 TGAAGTACAACTACTTGTGGGGAGGGAATGGCACAGAGCAAAAGATTGAAGGGAAGGAAG 3420  
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Db 3421 ATGGAGAGGCTCATNGTTTGGGGGTGTGAAGGTCACCTCTTTTCATGTGATGGAGGT 3480  
Qy 3481 TAAGAAAAACAGTGTGTAGTTTGTCTTTCAGACACACCCCAACTATCAAAACATATCC 3540  
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RESULT 3

US-09-354-243B-8  
; Sequence 8, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa  
; TITLE OF INVENTION: (TIFS)  
; FILE REFERENCE: LUD 5543.1  
; CURRENT APPLICATION NUMBER: US/09/354,243B



; CURRENT FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-354-243B-8

Query Match 100.0%; Score 7445; DB 3; Length 7445;  
Best Local Similarity 100.0%; Pred. No. 0;  
Matches 7445; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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QY	121	TAAGAGCGCTATATCAGCATTTAACCAACATGTTAATAGTTTCTTCTGCGCAAGCAACT	180
DB	121	TAAGAGCGCTATATCAGCATTTAACCAACATGTTAATAGTTTCTTCTGCGCAAGCAACT	180
QY	181	TGAATCTATGCTCTTAAACAACTTTCAAGCTCTTAATATAGTGTCAACGACTGGAGTCCG	240
DB	181	TGAATCTATGCTCTTAAACAACTTTCAAGCTCTTAATATAGTGTCAACGACTGGAGTCCG	240
QY	241	CTGCTGTCACACGAGCTCTTGACGACGCTCTCTCTGTTTGCAATTTTATGTTCTTTGA	300
DB	241	CTGCTGTCACACGAGCTCTTGACGACGCTCTCTCTGTTTGCAATTTTATGTTCTTTGA	300
QY	301	TCGACTCCCAACCTCTCACTTCCGCTCTGATGGCCACTTTCACATTTCTGCATTTA	360
DB	301	TCGACTCCCAACCTCTCACTTCCGCTCTGATGGCCACTTTCACATTTCTGCATTTA	360
QY	361	TGAATCTCATGTTTAACTCTTTTATTAATAATATTCACAAATCAGTGTGTCAGATC	420
DB	361	TGAATCTCATGTTTAACTCTTTTATTAATAATATTCACAAATCAGTGTGTCAGATC	420
QY	421	TGTTTCCACACATGATGCTGTGCACCAAGTCTGCTGCTGCTGCTGCTGCTGCTGCTG	480
DB	421	TGTTTCCACACATGATGCTGTGCACCAAGTCTGCTGCTGCTGCTGCTGCTGCTGCTG	480
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DB	601	TCTCTCTCCAGCCCAACATGAGTGTCTTTAGATTTCCACCTAGATAGATCTGATGGC	660
QY	661	TTCACTCAGTCGCACTCCCTTTGATCTTTTCTGCGCAAGGAAACCAAAAAAGCAAGAT	720
DB	661	TTCACTCAGTCGCACTCCCTTTGATCTTTTCTGCGCAAGGAAACCAAAAAAGCAAGAT	720
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DB	721	CCCACACTGCTTTCCGCTCTCAAGTCTGCACTCTCAACAGGTCAGATTTCTCCAGTGT	780
QY	781	CCCTCTAACACTTTCCCGAGTGCCTCTAACACTTTTCTCCAGTGTCCCTCTAACACTTT	840
DB	781	CCCTCTAACACTTTCCCGAGTGCCTCTAACACTTTTCTCCAGTGTCCCTCTAACACTTT	840
QY	841	CTCAGTGTCCCTCTAACACTTTTGTATCTCAATAGTGTAGGAGGAGAAAGATCTCACACA	900
DB			

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QY	901	GTGATTTTTCATGACTTTCGGCTTCTAGCTAGATGTAGGCAATTTGCGTGTGCTAGTCTAGG	960
DB	901	GTGATTTTTCATGACTTTCGGCTTCTAGCTAGATGTAGGCAATTTGCGTGTGCTAGTCTAGG	960
QY	961	AGGCGTGTGCTCCCGTGTAGGAAAGACTTTCCTAGTCTAGTGTGCTAGTGTGCTAGTGTG	1020
DB	961	AGGCGTGTGCTCCCGTGTAGGAAAGACTTTCCTAGTCTAGTGTGCTAGTGTGCTAGTGTG	1020
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DB	1021	GGATTCAGTGTACATCAATGCAAAAAATCCAGTATTTTGTAAATTTCTCTTCTTCAACT	1080
QY	1081	ATCCATCTATATATGTTTATTTAGTGTCTTAAATAATAATATTTTGGAGACTTATGC	1140
DB	1081	ATCCATCTATATATGTTTATTTAGTGTCTTAAATAATAATATTTTGGAGACTTATGC	1140
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DB	1141	TTGCAACAAGTAAATGTGTCAGAGAAATTTAGCAAAATGTATAGTATTTTATTTTAAAAAAA	1200
QY	1201	TCTATGCTTAAATGTCTATAGTGTTCACCTACCGATATTTCCAAACTTTAACTTGACC	1260
DB	1201	TCTATGCTTAAATGTCTATAGTGTTCACCTACCGATATTTCCAAACTTTAACTTGACC	1260
QY	1261	TTGGCTATGATTTTCAACCTTTGATTTTGATCTTACCATAACAGTCTCTGCAACCAAGACAT	1320
DB	1261	TTGGCTATGATTTTCAACCTTTGATTTTGATCTTACCATAACAGTCTCTGCAACCAAGACAT	1320
QY	1321	CTGTGGCAATGGGAGCTGTGAAAGAAAGCCAACTTCTTATTTAAAAAAAACAGCTA	1380
DB	1321	CTGTGGCAATGGGAGCTGTGAAAGAAAGCCAACTTCTTATTTAAAAAAAACAGCTA	1380
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DB	1381	GTATATAGTTTGTAGGATTTCCATATATAAATAAATAGAGATATATTTTAAAAAATGA	1440
QY	1441	AATAATCTCCAGTCTTTCATTTATGCTTATTTCAAAGCAGAGATATAGGACACGGGTCT	1500
DB	1441	AATAATCTCCAGTCTTTCATTTATGCTTATTTCAAAGCAGAGATATAGGACACGGGTCT	1500
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DB	1501	TTTATTTCTGGTCACTTCTAAAGAGATAAGAACTCTATGAAGTTGGTGGGAAATAGTCTC	1560
QY	1561	GTGACCAAAACGCTGACTCAATAGCTACCGGAGATCAAGGCTGCTCTACTCAATCAGAA	1620
DB	1561	GTGACCAAAACGCTGACTCAATAGCTACCGGAGATCAAGGCTGCTCTACTCAATCAGAA	1620
QY	1621	TCTACTACGGCAAGCCATGGCTTTCTTTGAAAAACGCTGTTTAAAGAGATTTCTGGGATTT	1680
DB	1621	TCTACTACGGCAAGCCATGGCTTTCTTTGAAAAACGCTGTTTAAAGAGATTTCTGGGATTT	1680
QY	1681	GTGTGCAAAAGCACTTGTGTGGCCCTCAACGTCGCTTTTGGGAAAGACTTCCCATCTCT	1740
DB	1681	GTGTGCAAAAGCACTTGTGTGGCCCTCAACGTCGCTTTTGGGAAAGACTTCCCATCTCT	1740
QY	1741	CAAGTGGGAAAGGCTTGGAGGTGCTGCTGTTGGCCCTCTTATGTTGGTGGTGGTACTTCTC	1800
DB	1741	CAAGTGGGAAAGGCTTGGAGGTGCTGCTGTTGGCCCTCTTATGTTGGTGGTGGTACTTCTC	1800
QY	1801	AGAAGACAGGACTTGGAAATTTAGATAATGCTGATGTCAATATCATTTACAAATACCAAAAA	1860
DB	1801	AGAAGACAGGACTTGGAAATTTAGATAATGCTGATGTCAATATCATTTACAAATACCAAAAA	1860
QY	1861	ACCCTGGTGTCCGATGGCTTATAAAAAGCAGCAACTTCTGCTCTCCCATCACAAGCAGAG	1920
DB	1861	ACCCTGGTGTCCGATGGCTTATAAAAAGCAGCAACTTCTGCTCTCCCATCACAAGCAGAG	1920
QY	1921	ACACCTTAAACAGGTAGGACTCAGACCTCTACAGCAATCATCTGCTTGGTACATGCTA	1980
DB	1921	ACACCTTAAACAGGTAGGACTCAGACCTCTACAGCAATCATCTGCTTGGTACATGCTA	1980

Qy	1981	CCGACGAA	CATGCTCC	CTGATGTT	TTTGGCTTT	TGCTCTCT	CACATAA	CAGGCTCT	CCCT	2040
Db	1981	CCGACGAA	CATGCTCC	CTGATGTT	TTTGGCTTT	TGCTCTCT	CACATAA	CAGGCTCT	CCCT	2040
Qy	2041	CTCACTTAT	CAACATGTT	TGACACAT	TGTGCGAT	CTCTGATG	GTGTCCT	CGCAGAAAT	CTATG	2100
Db	2041	CTCACTTAT	CNACTGTT	TGACACAT	TGTGCGAT	CTCTGATG	GTGTCCT	CGCAGAAAT	CTATG	2100
Qy	2101	AGTTTTTCC	CTTATG	GGGACCTT	TGGCCG	CCAGCTGC	CTGCTT	CTCATTTG	CCCTGTGGCC	2160
Db	2101	AGTTTTTCC	CTTATG	GGGACCTT	TGGCCG	CCAGCTGC	CTGCTT	CTCATTTG	CCCTGTGGCC	2160
Qy	2161	CAGGAGCA	AAATCGCT	CCCGCTCA	ACCCGCTG	CAAGCTTG	AGGTTG	CCAACTTCC	AG	2220
Db	2161	CAGGAGCA	AAATCGCT	CCCGCTCA	ACCCGCTG	CAAGCTTG	AGGTTG	CCAACTTCC	AG	2220
Qy	2221	CAGCCGTAC	ATCGTCAA	CCGACCTT	TATGTCG	CCCAAGAG	GTACAG	CTGCATCT	CTTT	2280
Db	2221	CAGCCGTAC	ATCGTCAA	CCGACCTT	TATGTCG	CCCAAGAG	GTACAG	CTGCATCT	CTTT	2280
Qy	2281	CTCTCCAT	ACCGCTT	GCACATTT	CTGAAAG	CACTTGC	AAACCTT	TAGGGCGC	TTTTA	2340
Db	2281	CTCTCCAT	ACCGCTT	GCACATTT	CTGAAAG	CACTTGC	AAACCTT	TAGGGCGC	TTTTA	2340
Qy	2341	TCCTCGAG	GTCTC	ACTACCTAT	GTCTCTCT	TTTAGAG	ACTCTT	TAAGGACT	CGGT	2400
Db	2341	TCCTCGAG	GTCTC	ACTACCTAT	GTCTCTCT	TTTAGAG	ACTCTT	TAAGGACT	CGGT	2400
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Db	2401	CTTTTCTAT	TCTATTT	CAAGGTCT	CAGGACCA	TTCCTAT	CTTTGG	CCCTTCAG	GACACA	2460
Qy	2461	TATACGAA	TTTTATCT	ACAGAGG	GCATTTAG	AAAGCC	ACCCAG	CTGCAAT	ACTTTC	2520
Db	2461	TATACGAA	TTTTATCT	ACAGAGG	GCATTTAG	AAAGCC	ACCCAG	CTGCAAT	ACTTTC	2520
Qy	2521	CATTCTCT	GTGCTCT	CTTGAACT	CACTCTCT	TTGGCTAC	CTCCTG	AGACCC	CACTGCG	2580
Db	2521	CATTCTCT	GTGCTCT	CTTGAACT	CACTCTCT	TTGGCTAC	CTCCTG	AGACCC	CACTGCG	2580
Qy	2581	GACATACAT	CTTACT	TACAGGCT	TTTTCTT	CCATCTCT	CTGAC	CCAGGCACT	TAGGGT	2640
Db	2581	GACATACAT	CTTACT	TACAGGCT	TTTTCTT	CCATCTCT	CTGAC	CCAGGCACT	TAGGGT	2640
Qy	2641	TTTCTCTCT	TTTACG	CCAGCTTGC	AGATAC	AAACAC	AGAGCT	CCGGCTCA	TCGGGGAGA	2700
Db	2641	TTTCTCTCT	TTTACG	CCAGCTTGC	AGATAC	AAACAC	AGAGCT	CCGGCTCA	TCGGGGAGA	2700
Qy	2701	AACTGTTCC	GAGGAGT	CACTGTA	GTGAGC	AGGCTAG	CTGCGGGAGC			2760
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Qy	2821	CTAAAGAT	CAGTCTA	CTGATGA	AGCAGGTG	CTCAACT	TCCCTG	TGAAAG	AGCTTCTGC	2880
Db	2821	CTAAAGAT	CAGTCTA	CTGATGA	AGCAGGTG	CTCAACT	TCCCTG	TGAAAG	AGCTTCTGC	2880
Qy	2881	TCCCCCAG	TCAGAC	AGTTCC	AGCCCTAC	ATGAGC	AGGTTG	TAATCT	TCTGACCCAAAC	2940
Db	2881	TCCCCCAG	TCAGAC	AGTTCC	AGCCCTAC	ATGAGC	AGGTTG	TAATCT	TCTGACCCAAAC	2940
Qy	2941	TCAGCAAT	CAGCTC	CTGTGT	TAAGTCT	GTACTCT	GTGCTAC	CTATGCT	CTCTCTCTT	3000
Db	2941	TCAGCAAT	CAGCTC	CTGTGT	TAAGTCT	GTACTCT	GTGCTAC	CTATGCT	CTCTCTCTT	3000
Qy	3001	CCTCTTCT	ATTTCA	GTAAAG	ACCCGAG	GTCTG	CCCTCT	CTCTCT	CTTCA	3060
Db	3001	CCTCTTCT	ATTTCA	GTAAAG	ACCCGAG	GTCTG	CCCTCT	CTCTCT	CTTCA	3060

Qy	3061	GGGCCTC	AGCAAC	CAACCAT	CATAGG	CCACTT	GAAATAG	GTCA	AAAGCCTT	TGGCTTC	3120
Db	3061	GGGCCTC	AGCAAC	CAACCAT	CATAGG	CCACTT	GAAATAG	GTCA	AAAGCCTT	TGGCTTC	3120
Qy	3121	AATTGAGT	ATAATCT	TGAGTTT	GTATG	AGTGAAG	CTTTAT	TTTGT	TATCCAT	TGGAAGA	3180
Db	3121	AATTGAGT	ATAATCT	TGAGTTT	GTATG	AGTGAAG	CTTTAT	TTTGT	TATCCAT	TGGAAGA	3180
Qy	3181	AATCAACT	CAAAATCT	TGTAGG	ATGAA	AGATGT	TGGGA	ACGAAA	AAAGCCT	TAGATAGA	3240
Db	3181	AATCAACT	CAAAATCT	TGTAGG	ATGAA	AGATGT	TGGGA	ACGAAA	AAAGCCT	TAGATAGA	3240
Qy	3241	GAACACAG	ATCTGCT	GTAGT	TAGTACT	TATATG	GGGGAG	CAGGGG	CGATAT	CCA	3300
Db	3241	GAACACAG	ATCTGCT	GTAGT	TAGTACT	TATATG	GGGGAG	CAGGGG	CGATAT	CCA	3300
Qy	3301	CAAGTACT	TTGTGG	GGAGAG	AAATCCA	CTGAGT	ACAAGT	ACTTGT	TGGCAT	TGGAGATCCAC	3360
Db	3301	CAAGTACT	TTGTGG	GGAGAG	AAATCCA	CTGAGT	ACAAGT	ACTTGT	TGGCAT	TGGAGATCCAC	3360
Qy	3361	TGAGTACA	AGTACT	TTGTGG	GGAGAG	AAATCCA	CTGAGT	ACAAGT	ACTTGT	TGGCAT	3420
Db	3361	TGAGTACA	AGTACT	TTGTGG	GGAGAG	AAATCCA	CTGAGT	ACAAGT	ACTTGT	TGGCAT	3420
Qy	3421	ATGGAGAG	GGCTCAT	TGTTGG	GGGTGT	GAAAGST	CACTCCT	TTTTCCA	TGTAT	TGAAGT	3480
Db	3421	ATGGAGAG	GGCTCAT	TGTTGG	GGGTGT	GAAAGST	CACTCCT	TTTTCCA	TGTAT	TGAAGT	3480
Qy	3481	TAAGAAAA	CAAGTGT	GTGAGT	TTGATG	CTTTCAG	ACACCC	CCCACT	TATG	AAACATATCC	3540
Db	3481	TAAGAAAA	CAAGTGT	GTGAGT	TTGATG	CTTTCAG	ACACCC	CCCACT	TATG	AAACATATCC	3540
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Db	3541	ACGAGGAG	CGGGAG	AGCTGT	GGGAG	ACCTG	GCATTTAG	GGAAG	CGCGCT	TTTCA	3600
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Db	3661	TTCTGTTT	CTACCG	TTCTTGT	CTACTG	TGTGGA	AACTTCA	GTAGG	ATTTCC	CAAGACGAGGA	3720
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Db	3721	CAGCTCTT	CTGTAA	GAGGAG	CCCTG	ATTTCA	GTGCTC	CTAGAG	AACTAG	CTCAGA	3780
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Db	3781	GAATCTAG	GTCAAC	CGTGAA	ATCTAG	GTCA	CGGGCA	AAAAAT	GACTGA	ACGCTCTATT	3840
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Db	3841	CCAGGTGAA	CCGTCA	CGTGCT	CAGATAT	ACTGAG	GTATTTGG	GTCTCC	ACCGGAT	TAGAT	3900
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Db	4021	AAGCAAT	TCAGT	AGGAG	CGTGGG	GAATTTCT	CTGCTT	CC	CAGTCC	CTTACTTTG	4080
Qy	4081	TAAATTTT	ATTTTCA	CTTGTCT	ACTTACT	CTGTCT	CAATTTAG	CTG	CACTGT	ATC	4140
Db	4081	TAAATTTT	ATTTTCA	CTTGTCT	ACTTACT	CTGTCT	CAATTTAG	CTG	CACTGT	ATC	4140
Qy	4141	TAGCTGG	GTCTAT	AGATCT	TTTCA	ATCTGT	GTCTTAA	ATTTGT	TAGT	CAAAATTTCTG	4200

Db 4141 ||||| TAGCTGGGCTCTATAGATCTTTCAATCTGTCTCTAAATTTGTAAGTCACAATTTCTGGAGCT 4200  
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Qy 7441 TGTAT 7445  
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RESULT 4

US-09-178-973B-17  
; Sequence 17, Application US/09178973B  
; Patent No. 6274710  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFs)  
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543  
; CURRENT APPLICATION NUMBER: US/09/178,973B  
; CURRENT FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 17  
; SEQ ID NO 17  
; LENGTH: 5935  
; TYPE: DNA  
; ORGANISM: Mus musculus  
US-09-178-973B-17

Query Match 57.0%; Score 4245.2; DB 3; Length 5935;  
Best Local Similarity 88.3%; Pred. No. 0;  
Matches 5039; Conservative 0; Mismatches 178; Indels 487; Gaps 20;

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Qy 293 TACCATGCTATCCGACGAGCATGTTCCCTCGATGTTTTCGCTTTTTCGCTCTCTCCTCACTAAC 352  
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3583 GGCGCGCTTTTCAACGAGAAATTTATGCTCATCTCTTGTGCTACACTCCACCTTTG 3642  
Db GGCGCGCTTTTCAACGAGAAATTTATGCTCATCTCTTGTGCTACACTCCACCTTTG 1915  
3643 ATGAGGTTACGCTCAGGTTTCTGTTTCTACCGTCTTCTGCTACTGTTGGAACCTTCACTAGG 3702  
Db ATGAGGTTAAAGCTCAGGTTTCTGTTTCTACCGTCTTCTGCTACTGTTGGAACCTTCACTAGG 1975

3703 ATTCCCAAAGACGAGGACAGCTCTTCTGTAAAGGAGGACCTGGATTTTCAGTGTCTAG 3762  
Db ATTCCCAAAGACGAGGACAGCTCTTCTGTAAAGGAGGACCTGGATTTTCAGTGTCTAG 2035  
3763 AGAACGAATAGCTCAGAGATCTAGGTCAACCTGAAATCTAGGTCAACAGGGGCAAAA 3822  
Db AGAACGAATAGCTCAGAGATCTAGGTCAACCTGAAATCTAGGTCAACAGGGGCAAAA 2095  
3823 TGAAGTCAACGCTTATTTCCAGGTGAACGCTCAGGTCACTGACATATCTAGGTATGG 3882  
Db TGAAGTCAACGCTTATTTCCAGGTGAACGCTCAGGTCACTGACATATCTAGGTATGG 2155  
3883 GCTCCACCGGATTAAGATTCTGTAGTCTGCTTTTATTTTTCAGCAGCATCAGCGGT 3942  
Db GCTCCACCGGATTAAGATTCTGTAGTCTGCTTTTATTTTTCAGCAGCATCAGGTGT 2215  
3943 GACGACGAGCAATCCAGAAAGATGTCAGAGGCTGAGGAGACAGTGAAGAAAGTACTA 4002  
Db GACGACGAGCAATCCAGAAAGATGTCAGAGGCTGAGGAGACAGTGAAGAAAGTACTA 2275  
4003 TTGGCAAGCCACAATTAAGCCATTTAGTAGAGAGCTGGGATTTCTTCTCTGCTTC 4062  
Db TTGGCAAGCCACAATTAAGCCATTTAGTAGAGAGCTGGGATTTCTTCTCTGCTTC 2335  
4063 CCAGTCCCTTCTACTTTGTAAATTTTATTTGACTTGTCTATCTATCTGCTGCTTCTCG 4122  
Db CCAGTCCCTTCTACTTTGTAAATTTTATTTGACTTGTCTATCTATCTGCTGCTTCTCG 2395  
4123 CTTAGCTGCACCTGTATCTAGCTGGGTCTATAGATCTTTCAATCTGTGCTTCAATTTGTA 4182  
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4183 AGTCACAAATTTCTGAGCTAGCAAGAGCTTAGCTCAGCCAGTCTCATGAGCACTTGTCTCG 4242  
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4243 GAGGATGGCTGTGACAGAGTCAATGCTAGAAAGAGCATCCCTGATTTCCAGCTCTGCA 4302  
Db GAGGATGGCTGTGACAGAGTCAATGCTAGAAAGAGCATCCCTGATTTCCAGCTCTGCA 2575  
4303 CTTGCTTAGTGGCCATGTGTAAATTTACTTTGGCTTGTATTAAGTATTTGGGAAAGCAGTTC 4362  
Db CTTGCTTAGTGGCCATGTGTAAATTTACTTTAGCTTGTAAAGTATTTGGGAAAGCAGTTC 2635  
4363 CCACGACCTTACATAATCTGAAAGACCATGTCATTTGAAAGCTAGAAAGCTGGGCAAACT 4422  
Db CCACGACCTTACATAATCTGAAAGACCATGTCATTTGAAAGCTAGAAAGCTGGGCAAACT 2695  
4423 TACTAGAGATGATTTTGTAGCTCATTTAAAGCGATGCTCTGAAATGTGGCAAAATCAACCC 4482  
Db TACTAGAGATGATTTTGTAGCTCATTTAAAGCTCATTTAAAGCTGCTGAAATGTGGCAACCC 2755  
4483 AGAATAACAAAGAGCTGGATTTGCAAAATAGGACAAGTATTTAGAATCACTGGTATT 4542  
Db AGAATAACAAAGAGCTGGATTTGCAAAATAGGACAAGTATTTAGAATCACTGGTATT 2815  
4543 AATAGCTATCATCTTAATTAATAATAGGCGCTATATA---TATATTTAAGATTAACA 4598  
Db AATAGCTATCATCTTAATTAATAATAGGCGCTATATA---TATATTTAAGATTAACA 2875  
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4719 TCAATTTCTTAGT-----TTTTTTTTTCTTCAACCCCTGATCAAGCCACTAGTAGGACCT 4773  
Db TCAATTTCTTAGT-----TTTTTTTTTCTTCAACCCCTGATCAAGCCACTAGTAGGACCT 3055  
4774 ATCTGCTGAGCTATATATGACTTTTACAGCAACAAACATTTCTGCTGTGGCCTCTTTGG 4833

||||| 3056 ATCTGCTGCGAGCTATTATATGACTTTACAGCAACCAACATTGCTGTGTGCGCTCTTTGG 3115  
QY  
4834 GGAAGGGAACAGGATAGCAGAGAGGCTCAGGCTAGCAAGTCT-GACTTGGCCCTAAAGCCAG 4892  
Db  
3116 GGAAGGGAACAGGATAGCAGAGAGGCTCAGGCTAGCAAGTCTGGACTCAACCTAAAGCCAG 3175  
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Db  
3176 AGGCATGGTGTATAGCAGAGAAAGTGAGGCTCTTCACAAAGTGGGTGCTTAAGTAATCA 3235  
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4953 GAAACAGGAAGGCTCCGGTGTGATGGAATTAATCAGTAAGATATCTACCCCTTATCTC---CT 5009  
Db  
3236 GAAACAGGAAGGCTCTGGTGTGATGGAATTAATCAGTAAGATATCTACCCCTTATCTCTTCT 3295  
QY  
5010 TCTATCGAACCCTAAATCGTCTCTTTTCTGTGTGTGTAGGCTGATAAACAACACTTGTTTTC 5069  
Db  
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Db  
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QY  
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Db  
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QY  
5610 TCTGGAGGCGAGCAAG--TGAGGGAATGGAAGGGAAGGAAGGAAGATGTAGAGGA 5666  
Db  
3895 TCTGGAGGCGAGCAAGCAITTGAGAGGAGATGGAAGGGAAGGAAGGAAG--AATGTAGAGGA 3952  
QY  
5667 CTTGAAACAGCTACAAATCCTCTACACAGACGATTTTCTTGTGAAACAATCTAGAA---GGT 5722  
Db  
3953 TTTGAAAGCTACAAATCCTCCACAGAGGATTTTCTTGGAGGAATCTAGAACAGGGT 4012  
QY  
5723 AGTGGAATAGGT-GAATCGAGGGGACTTGTCTTGGCAATTTGAATCTGGGTTTTTGTCTC 5781  
Db  
4013 GGTGAATTAGGTGGATCGCAGAGAGCTTGTCTTGGCAATTTGAATCTGGGTTTTTGTCTC 4072  
QY  
5782 TCCATTGAGTGAAGCGTCAACC-TTTTTCCTCGAATGGAGGGAAGAGGGGT 5840  
Db  
4073 TCCATTGAGTGAAGCGTCAACCCTTTTACCTCGAATGGAGGGAAGAGGGGT 4132  
QY  
5841 GTTATGACTCTCTACCTCGAGTGTACTAGTTTACGCAATGGAAACAGACACTCGGGACCTC 5900  
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Db  
4133 GTTTTGACTCTACTCGAGTGTCTTAGTGTATTAGCAATGGAACAGACACTCGGAGCCTC 4192  
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5901 CTCTTGAC-----AAAAAAATGGAAACCTGTTGTTTGTCTTGTGTTTGTCTTTTG 5950  
Db  
4193 CTCTTGACAAAGAAAAAAGGAAACCTGTTGTTTCTCTTGTGTTTGTCTTTTG 4252  
QY  
5951 TTAAAGAAAGCAC----- 5963  
Db  
4253 TTAAAGAAAGCACAGGCAGCTGGGCCATGCTGTCCTTTTAAATCCAGCAITTTGGAG 4312  
QY  
5964 ----- 5963  
Db  
4313 GCAGAGCAGGTGACTTTCTTAAATTTCAAGGCCAGCCTGCTTACAAAGTGAGTTCAGGA 4372  
QY  
5964 ----- 5963  
Db  
4373 CAGCCAGGCTATACAGAGAAACCTGCTCTCGGGAATAAAAAAAGAAAGAAAG 4432  
QY  
5964 ----- 5963  
Db  
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QY  
5964 ----- 5963  
Db  
4493 GAGAGGAGAGGAA 4552  
QY  
5964 ----- 5963  
Db  
4553 GAGAAGAA 4612  
QY  
5964 -----GGCAAGCCCGACCACATCGGT 5985  
Db  
4613 AAGAGAAAGAAAGAAAGCAAGCAAGCACTGGCAAGCATGCCACATGGGA 4672  
QY  
5986 TGAATGTGGTCTTTGAGTCAAGGCTTTTGTGAGTGAACAATCAATAGTTGATCGG 6045  
Db  
4673 CGTATGTGGTCTTTGAGCAAGGCTTTTGAATTTGAGCGCTCATCAATAGTTGATCGG 4732  
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6046 TCAGGTGGAGGCTTACCTGTTCAGCGCGCCCTGCTGGCTTCGCACCTTACATCTCCAGG 6105  
Db  
4733 TCAGGTGGAGGCTTACCTGTTCAGCGCGCCCTGCTGGCTTTAGCATTACATCTCCAGG 4792  
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Db  
4793 TCTCAGTATCACTTCTCTGCTGCTTTAGCACAGTTAGGAGTTGAGCAAACTTTTTCCTCA 4852  
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Db  
4853 CCCCCACTAAATTTAAATTTGACAAAAGCTGTGTAATTTGTGGGATACAGTGTGATAAT 4912  
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Db  
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QY  
6286 TATGGGTGTGAAATGCAAGTAATATAGGTAGATCCCTGTGTGTGCTTCTTAGGTTCAGAAAG 6345  
Db  
4973 TATGGGTGTGAAATGCAAGTAATATAGGTAGATCCCTGTGTGTGCTTCTTAGGTTCAGAAAG 5031  
QY  
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Db  
5032 CATGATTTTAAAGTCTTTGGGCAAAATCATATATCTCATGCTTAAATATCATTTATGTGA 5091  
QY  
6406 TTATTATCTTTTAGAAGAGCTGATCTTGGTTTTTGGTGTCTCAGCAGCAAAATGTCAAC 6465  
Db  
5092 TTATCAATCTTTTAGAAGAGCTGATCTTGGTTTTTGGTGTCTCAGCAGCAAAATGTCAAC 5151  
QY  
6466 AGCTCTTTTAACTGGTACCACTTTAGAAAATGCTACCTGTGCTCAAAATTTGGTTGTATT 6525  
Db  
5152 AGCTCTTTTAACTGGTACCACTTTAGAAAATGCTACCTGTGCTCAAAATTTGGTTGTATT 5211  
QY  
6526 CTTATTTTCAATAGCTTCGAGAGAGTGGAGAGATCAAGCGGATTCGCGGAACTCGGACCTGCT 6585  
Db  
5212 CTTATTTTCAATAGCTTCGAGAGAGTGGAGAGATCAAAAGCGATCGGGAACTCGGACCTGCT 5271  
|||||





Qy	2991	CCTCTCTCTTCCCTCTTCTATTTCCAGTAAGAACCGAGGCTCTGCGCTCTCTCTCTTCCACA	3050
Db	1312	CCTCTCTCTTCCCTCTTCTATTTCCAGTAAGAACCGAGGCTCTGCGCTCTCTCTCTTCCACA	1371
Qy	3051	AGAGTGAGGAGGGGCTCAGACACACACCACTCATPAGGCCACTTGAATAGGTCACAAGG	3110
Db	1372	AGAGTGAGGAGGGGCTCAGACACACACCACTCATPAGGCCACTTGAATAGGTCACAAGG	1431
Qy	3111	CTTTGGCTTCAATTGAGTAATACTTTGAGTTTGTATGAGTGAAGCTTTATTTGTTTATC	3170
Db	1432	CTTTGGCTTCAATTGAGTAATACTTTGAGTTTGTATGAGTGAAGCTTTATTTGTTTATC	1491
Qy	3171	CATGGAAGAAATCAACTCAAAATTTCTGATGATCAGAAAGATGTTGGGAACGAAAAAGG	3230
Db	1492	CATGGAAGAAATCAACTCAAAATTTCTGATGATCAGAAAGATGTTGGGAACGAAAAAGG	1551
Qy	3231	CCTAGATAGAAAAACAGATCTGCTGAGTATAGTACTTAT ---GGGGGGAGCAGGGGGCG	3286
Db	1552	CCTAGATAGAAAAACAGATCTGCTGAGTACAGTACTTATGGGGGGGGGGCAGGGGGCG	1611
Qy	3287	ATATCCACTGAGTACAAGTACTTGTGGGGAGAGAAATCCCACTGAGTACAAGTACTTGTG	3346
Db	1612	ATATCCACTGAGTCCAAGTACTTGTGGGGAGAGAAATCCCACTGAGTACAAGTACTTGT--	1689
Qy	3347	GCATGGAGATCCACTGAGTACAAGTACTTGTGGGGGGAGGGAATGGCACAGAGCAAAAGT	3406
Db	1670	-----GGGGGAAGGATGGCACAGAGCAAAAGT	1697
Qy	3407	TGAAGGGA---AGGAAGATGAGAGGGCTCATGTTGGGGGTGTGAAGGTCACTCC--TT	3462
Db	1698	TGAAGGGAAGAGAGAGATGGAGGGCTCAATGTTGGGGGTGTGAAGGTCACTCCCTTT	1757
Qy	3463	TTCCATGTGATGGAGAGTTAAGAAAAACCAAGTGTGTGAGTTTGATGTCTTCAGACACCCC	3522
Db	1758	TTCCATGTGATGGAGAGTTAAGAAAAATCAGTGTGTGAGTTTGATGTCTTCAGACACCCC	1817
Qy	3523	CAACTATGAACATATCCACGAGGAGCGGGCAGACTGTGGGAGACCTGGCAATTTAGGGAA	3582
Db	1818	AA-----CTATGGCAGACTGTGGGAGACCTGGCAATTTAGGGAA	1855
Qy	3583	GGCGGGCTTTTCACGAGAAACTTTATGCTCATCTCTGTGCTACACTCCCACTTTG	3642
Db	1856	GGCGGGCTTTTCACGAGAAACTTTATGCTCATCTCTGTGCTACACTCCCACTTTG	1915
Qy	3643	ATGAGGTTCACTCAGGTTTGGTTTCTACCGTTCTGTCTACTGTGGAACTTCAGTAGG	3702
Db	1916	ATGAGGTTAAGCTCAGGTTTGGTTTCTACCGTTCTGTCTACTGTGGAACTTCAGTAGG	1975
Qy	3703	ATTCCTCCAAAGCAGGACAGCTCTTCGTAAAGGGGAGCTCGAATTCAGTGTCTTAG	3762
Db	1976	ATTCCTCCAAAGCAGGACAGCTCTTCGTAAAGGGGAGCTCGAATTCAGTGTCTTAG	2035
Qy	3763	AGAACGAAATAGCTCAGAGAATCTAGGTCACACGTGAATCTAGGTCAAGCGGGCAAAA	3822
Db	2036	AGAACGAAATAGCTCAGAGAATCTAGGTCACACGTGAATCTAGGTCAAGCGGGCAAAA	2095
Qy	3823	TGACTGAACGGCTTAATTCAGAGTGAACGGTCACTGCTCAGATATCTGAGGTATTGG	3882
Db	2096	TGACTGAACGGCTTAATTCAGAGTGAACGGTCACTGCTCAGATATCTGAGGTATTGG	2155
Qy	3883	GCTCCACCGGATAGATTTCTGTAGTCAAGTCTGCTTTATTTTGCAGCATCAGCGGT	3942
Db	2156	GCTCCACCGGATAGATTTCTGTAGTCAAGTCTGCTTTATTTTGCAGCATCAGTGGT	2215
Qy	3943	GACGACGAGAACATCCAGAGAATGTCAAGAGGCTGAAGGAGACAGTGAAGAGGTACTA	4002
Db	2216	GACGACGAGAACATCCAGAGNATGTCAAGAGGCTGAAGGAGACAGTGAAGAGGTACTA	2275
Qy	4003	TTGGCAAGCCCAATPACTAAGCCATTCAGTAGGAGACGTGGGATTTCTTCTCTGCTTC	4062
Db	2276	TTGGCAAGCCCAATPACTAAGCCATTCAGTAGGAGACGTGGGATTTCTTCTCTGCTTC	2335

Qy	4063	CCAGTCCCTTCTACTTTGTAAACATTTTATTTGACTTGTCTACTATCTGTCCTCAATTACTCG	4122
Db	2336	CCAGTCTCTTCTACTTTGTAAACATTTTCTTTGACTTGTCTACTATCTGTCCTCAATTACTCA	2395
Qy	4123	CTTAGCTGCACCTGTATCTAGCTGGGTCTATAGATCTTTCAATCTGTGTCTAAATTTGTA	4182
Db	2396	CTTAGCTGCACCTGTATCTAGCTGGGTCTATAGATCTTTCAATCTGTGTCTAAATTTGTA	2455
Qy	4183	AGTCACAAATTTCTGAGCTAGCAGAAAGCTTTAGCTCAGCCAGTCTCATGAGCAGCTTGCTCG	4242
Db	2456	AGTCACAAATTTCTGAGCTAGCAGAAAGCTTTAGCTCAGCCAGTCTCATGAGCAGCTTGCTCG	2515
Qy	4243	GAGGATGGCTTTGTGACAGAGTCAATGCTAGAAGACAGCATCCCTGATTTCCAGCTCTGCA	4302
Db	2516	GAGGATGGCTTTGTGACAGAGTCAATGCTAGAAGACAGCATCCCTGATTTCCAGCTCTGCA	2575
Qy	4303	CTTGGCTTAGTGGCCATGTGTAAATTACTTTGGCTTGTATTAAGTATTTGGGAAAGCCAGTTC	4362
Db	2576	CTTGGCTTAGTGGCCATGTGTAAATTACTTTAGCCCTGATTAAGTATTTGGGAAAGCCAAATTC	2635
Qy	4363	CCACGGACCTACATAATCTCAAGAACCATGCTATTGAAAACTAGAAAAGCTGGGCACAAACT	4422
Db	2636	CCACGGACCTACATAATCCGAAAGCATGCTATTGAAAACTAGAAAGCTGGGCACAAACT	2695
Qy	4423	TACTAGAGATGATTTTGTAGCTCATTTAAACCGGATGCTCTGAAATGTGGCAAAATCAACCC	4482
Db	2696	TACTAGAGATGATTTTGTAGCTCATTTAAACCTGATGCTCTGAAATGTGATCAAAATCAACCC	2755
Qy	4483	AGAATAACAACAAAGAGCTGGATTTGCAAAATAGGACAGATTTTAGAATCACTGGTATT	4542
Db	2756	AGAATAACAACAAAGAGCTGGATTTGCAAAATAGGACAGATTTTAGAATCACTGGTATT	2815
Qy	4543	AATAGCTATCATCTTAATTAATAATATAGGCGCTATATA ---TATATTAAGATTAACA	4598
Db	2816	AACAGCTGTCACTTAATTAATAATATAGTGTCTATTAGTGGCTATTTTAGATTAACA	2875
Qy	4599	CAAGAGTGGATAGCCTCCCAATTTTACTTGGCTGGTTTCAAAAAGAGTAAATAATCAGTC	4658
Db	2876	CAAGAGTGGATAGCTTCCCAATTTTACTGGCGCTGGTTTCAATAGAGTAAATAATCAGTC	2935
Qy	4659	ATGATTAATTAATAGTGTGCATGAAGATATGAGATGGAAACCCCTTCCCTTACTTTTACCT	4718
Db	2936	ATGATTAATTAATAGTGTGCATGAAGATATGAGTGGAAACCCCTTCCCTTACTTTTACCT	2995
Qy	4719	TCATTTCTTAGT ---TTTTTTTTTCTTCACACCCCTGATCAAGCCACTAGTAAGCACCT	4773
Db	2996	TCATTTCTTAGTATTATTTTTTTTTTCTTCACACCCCTGATCAAGCCACTAGTAAGCACCT	3055
Qy	4774	ATCTGCTGTGAGCTATTATATGACTTTTACAGCAAAACAACTTGTGTGGCCCTCTTTGG	4833
Db	3056	ATCTGCTGTGAGCTATTATATGACTTTTACAGCAAAACAACTTGTGTGGCCCTCTTTGG	3115
Qy	4834	GGAAGGAAACAGGATAGCAGGAGCTCAGGCTAGCAAGTCT- GACTTGGCCCTAAGCCAG	4892
Db	3116	GGAAGGAAACAGGATAGCAGGAGCTCAGGCTAGCAAGTCTGGAACCTCAACCTAAGCCAG	3175
Qy	4893	AGGCATGTTGATAGCAGAGAAAGTGAGGCTCTTCGCAAGTGGGTGTCTTAAAGTAAATCA	4952
Db	3176	AGGCATGTTGATAGCAGAGAAAGTGAGGCTCTTCACAAAGTGGGTGTCTTAAAGTAAATCA	3235
Qy	4953	GAACACAGAAAGGCTCCCGTTGATGGAATTAATCAGTAAGATATCTACCTTATCTC ---CT	5009
Db	3236	GAACACAGAAAGGCTCTCGTTGATGGAATTAATCAGTAAGATATCTACCTTATCTCCTTCT	3295
Qy	5010	TCATATCGAAACCTAAATCGTCTCTTTTCTGTGTGTAGGCTGATAAACAACACTTGTTTTC	5069
Db	3296	TCATATAGAACTAAACCGCTCTCTCCTTTCTGTGTGTAGGCTGATAAACAACACTTGTTTTC	3355
Qy	5070	TTTTGAGTGTTCATGGCTTTGTAGATGCTCTGCCAGTCTCTGTGTAGAGGGTTT	5129
Db	3356	TTTTGAGTGTTCATGGCTTTGTAGATGCTCTGCCAGTCTCTGTGTAGAGGGTTT	3415
Qy	5130	GTTACCTTGACACCTGGGCTTTGGATGTTAGCATGCCAAAGGCACACACTTCTGAATGCCT	5189



Db 3416 GTTACCTTGACCTGGCTGGATGTAGCATGCCAAAGGCGACACACTTCTGAATGCCT 3475  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5190 GTGTAAAGGTTATTATTCTATTACTTGTCTTTGGAAAGGTGAAGCGTGTGTGAGAAAG 5249  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 3476 GTGTAAAGGTTATTATTCTATTACTTGTCTTTGGAAAGGTGAAGTGTGTGAGAAAG 3535  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5250 AACTCAGAGAGATGTCTCTGTAGTAAAGTCTTTTCTTTTCCCTTAAATCCCTATAA 5309  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 3536 AACTCAGAGAGATGTATTCTGTAGTAAAGTCTTTTCTTTTCCCTTAAAGGCTATAA 3594  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5310 TCCACTTTGAGTCAACTTTGACCTTTTATACATGCTGTACATGAAGAGTGTTAGGCC 5369  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 3595 TCCACTTTGAGTCAACTTTTATACATGCTGTACATGAAGAGTGTTAGGCC 3654  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5370 CGCTCTCATGGCTCTGGGAAAGCACCATAAGGGAAGGAATGTTATGCTGAGAAATCTG 5429  
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Db 3655 CGCTCTCGTCTCTGGGAAAGCACCATAAGGGAAGGAATGTTATGCCGAGAAATCTG 3714  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5430 ACCGGCAGGGAACCTGGTCAAGAGCTCCCGGAAGACACACAGGTGTTAAGTAGGAACA 5489  
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Db 3715 ACTGGCAGGGAACCTGGTCAAGAGCTCCCGGAAGACACACAGGTGTTAAGTAGGAACA 3774  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5490 GTCAGGCTGGCTCATGTATAGAAATGGAACAGAGCGGGAAGATAGCTCAAGTT 5549  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 3775 GTCAGGCTGGGTTCATATAATAGAAATGGAACAGAGCGGGAAGATAGCTCAAGTT 3834  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5550 TCATAGGCTCGGAGTCTTAAAGATACAAATAGCTCTTGGGCTTCATACCAAGGAAG 5609  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 3835 TCATAGGCTCTTAAGTCTTTAAAGATACAAATAGCTCTTGGGCTTCATACCAAGGAAG 3894  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5610 TCTGGGAAGCAGCAAG- --TGAGAGGGAATGGAAGGGAAGGAACAGAAATGTAGAGGA 5666  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 3895 TCTGGGAAGCAGCAAGCATTGAGAGGAGATGGAAGGGAAGGAAC- --AATGTAGAGGA 3952  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
Db 5667 CTTGAAACAGCTACAAATCTCTACAGACAGATTTTCTTGGAACAATCTAGAA- --GGT 5722  
Qy ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
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RESULT 6
US-09-354-243B-29
; Sequence 29, Application US/09354243B
; Patent No. 6359117
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louned, Jamila
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa
; TITLE OF INVENTION: (Title)
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.1
; CURRENT APPLICATION NUMBER: US/09/354,243B
; CURRENT FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 29
; LENGTH: 5935
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-09-354-243B-29

Query Match 57.0%; Score 4245.2; DB 3; Length 5935;
Best Local Similarity 88.3%; Pred. No. 0;
Matches 5039; Conservative 0; Mismatches 178; Indels 487; Gaps 20;

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Qy 5986 TGAATGTGGGCTTTTGAAGTCAAGGCTTTTGAAGTTGAGCACTCATCAATAGTTGATCATGG 6045  
Db 4673 CGTATGTGGGCTTTTGAAGCAAGGCTTTTGAATTTGAGCGCTCATCAATAGTTGATCATGG 4732  
Qy 6046 TCAGGTGAGGGCTACCTGTGAGGCGAGCCCTGCTGGCTTCGCACTTAAACATCTCCAGG 6105  
Db 4733 TCAGGTGAGGGCTACCTGTGAGGCGAGCCCTGCTGGCTTTAGCACTTAAACATCTCCAGG 4792  
Qy 6106 TCTCAGTATCACTTCTCTACTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTCCAA 6165  
Db 4793 TCTCAGTATCACTTCTCTCTGCTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTCCAA 4852  
Qy 6166 CCCCACATAAATTTAATTTGACAAAGACTGTGTAATTTTGGGATACAGTGTGATTAAT 6225  
Db 4853 CCCCACATAAATTTAATTTGACAAAGCAGTGTAATTTTGGGATACAGTGTGATTAAT 4912

Qy 6226 GATCTATGTGTGATTTGTGCAAGGTTCAATAAGATAGATTAAATAGGCCCATCAACAGCTT 6285  
Db 4913 GATCTATGTGTGATTTGTGCAAGGTTCAATAAGGTAGATCAATAGGCCCATCAACAGCTT 4972  
Qy 6286 TATGGGTGTAATGCAAGTAATATAGGTAGATGCCCTGTGGTGTCTTCTTAGGTGAGAAAG 6345  
Db 4973 TATGGGTGTAATGCAAGTAATATAGGTAGATGCCCTGT--GTGCTCTTAGGTGAGAAAG 5031  
Qy 6346 CATGATTTTAAGTCTTTGGGCAATCATATTATCTCATGCTAAATAATACATTATGTTGA 6405  
Db 5032 CATGATTTTAAGTCTTTGGGCAATCATATTATCTCATGCTAAATAATGCAATTATGTTGA 5091  
Qy 6406 TTATTAATCTTTTAGAAGAGGCTGATCTTGGTTTTGGTGTCTCAGCAAGCAAAATGTCAAC 6465  
Db 5092 TTATCAATCTTTTAGAAGAGGCTGATCTTGGTTTTGGTGTCTCAGCAAGCAAAATGTCAAC 5151  
Qy 6466 AGCTCTTTTAACTGGTACCACTTTAGAAAATGCTACCTGTGTCTCAAAATTTGGTTGTATT 6525  
Db 5152 AGCTCTTTTAACTAGTACCACTTTAGAAAATGCTACCTGTGTCTCAAAATTTGGTTGTATT 5211  
Qy 6526 CTTATTTTCACTAGCTTTGGAGAGATCAAGGCGATTTGGGGAACCTGGACCTGCT 6585  
Db 5212 CTTATTTTCACTAGCTTTGGAGAGCGAGAGATCAAGGCGATCGGGGAACCTGGACCTGCT 5271  
Qy 6586 GTTTATGTCTCTCAGAAAATGCTTGGCTCTGAGCGAGAAAGCTAGAAAAACGAAGAACTG 6645  
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Qy 6706 AAGTGAGAGCTAACGTCCTCATCATTTAGAAAGTTTCAATGAAACCTGGCTCAGTTGA 6765  
Db 5392 AAGTGAGAGCTAACGTCCTCATCATTTAGAAAGTTTCAATGAAACCTGGCTCAGTTGA 5451  
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Qy 7006 AGTTTATTAATATGGAATTTTATATAGAAAACATTTATCTGCTATTTGATATTT-AGTATAAG 7064  
Db 5692 AGTTTATTAATATGGAATTTTATATAGAAAATTTATCTGATGTTGATATTTAGTATAAA 5751  
Qy 7065 GCAAAATATTTATGCAATTAATGGAACAGAGATATCTTAGGCTTTTAATAAACAACA 7124  
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Db 5812 TGAATATCATAAATCTTCTGTCTTGTAAATTTTCTCCCTTTAATATCAACNATACCATCA 5871  
Qy 7185 TCATCATCATTTACCCTTCTCATGATTTTCAATGCTTGAAGTATTTATGCTGTTTAA 7244  
Db 5872 TCGTCATCATTTACCCTTCTCATGACTTTCATGCTTGACTCATATATTTCTGTAAG 5931  
Qy 7245 GTTG 7248  
Db 5932 TTG 5935







Db 4940 TGAAGAAACCA--ACAATGAAAGCTTGTAGACTAACTAATAGTACTACACCCCAAAACCGGA 4997  
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Db 4998 GGAATGATTAGGAGCAGTGAAGTGTAGGCTCTT--GCAAGCAGGTACAACTAAATACCTCAG 5056  
Qy 4954 AAACAGGAAGGCTCCGGTGTGATGGAATATACGTAAGA-----TATCTACCCCTTA 5003  
Db 5057 AAACATGAAAGGCTCCAGTGTGATGGAATTTTTCAGTAAACAAGCTTAAACCTTAATTTCCCCCTT 5116  
Qy 5004 TCTCCCTCTATCGAACTTAATCGTCTCTTTCTTCTGTGTGTAGGCTGATAAACACACTT 5063  
Db 5117 TTTCCCTCTTGACTTTTAAAGAGGCTTTCTTCCGAGCATCATTTAATGAGTGTGACT 5176  
Qy 5064 GTT--TTCTTTTGTAGTGTTCATGGCTTTGTAGATTTTGTAGTGTCTGCGCAAGTTCCTGT-- 5119  
Db 5177 GTTCTCTCTTGTGATAATTGAAGCTTTGTAGTTTAAATTTGTGAAGCCCAAGTTCCTTGTG 5236  
Qy 5120 TAGAGGTTTGTACCTTGTACACCTGGGCTTTGGATTTAGATGTAGCATGCGAAAGGCACACACTT 5179  
Db 5237 TTATAGAACTATTATCTAGACATGGAGGCTGAATGTTAGCATGCGCCACAGACAAAGGCATG 5296  
Qy 5180 CTGAATGCTGTGTAAAGGTTATTATTCAATTTACT-----TTGCTCTTGGAAAGGT 5231  
Db 5297 CTTTACACATCTTGTCTTAAAGAAATTTACTGATTTCACTTGTCTGTGTCTTTTGAAGAAAGT 5356  
Qy 5232 GAAGCGTGTGTGAAGAAAGAACTCACAGGAGATGTCTCTGTAGGAAAGAACTTTTCTTTT 5291  
Db 5357 GAAGTGTGAGAGAGGAAATCTCATGTG-----ATCTGTGT 5393  
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Qy 5349 ACATGAAGAGTGTTAGGCCCCCTCTCATGGCTCTGGGAAAGACCAATAGGGAGG 5408  
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Qy 5463 -----GACCACCAACAGGTGTTAAGTAGGAACAGTCCAGGGTGGGCTCAT 5506  
Db 5573 GGTTTTGGGAGGCATTAAATCCCTCTCTGTTGGGGTAAAGAGCAGAACCGCAGGTGGTGTAGT 5632  
Qy 5507 GTAATAGAAATGGAACAGAGCGAGGAAGATAGCTACAAAGTTTCATAGGCTCC-GGAGT 5565  
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Qy 5624 AAGTGAAGGGAATGGAAGAGGGAAGAAACAGAAATGTAGAGGACTTTGAACAGCTTACAAAT 5683  
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Db 5800 CAATGGTGACAGGTTTTTCTTGAATTAATGCAAAATATGATAGATAGAGGAAATTTTCAGT 5859  
Qy 5744 GGGACTTGTCTTGCATTTGCAATCTGGGTTTTTGTCTCTCCATTTAGGTTGAAAGCGTCA 5803  
Db 5860 AGGGAATGCTTTTCACTTGAATTTGGGTTTTCTCTCT----TCGATTAAGTTTGGGATCTCTCA 5916  
Qy 5804 CCCTTTTACCTTCGAATGGAGGAGGAAGAGGGGTGTTATGACTCCTTACCTCGGAGTTT 5863  
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Qy 5864 TACTAGTTTACGCAATCGAACACAGACACTCGGGACCTCTCTTGTGACAAAGAAATGGAAC 5923  
Db 5973 TATTAACTAAAGCAAGTGGAAAGAGACTTATTGTGATTTTCTCCACAAAGAGTGAAAGACTT 6032

Qy 5924 CTGTCTTTTGTCTTGTGTTGTTCTTTTGTAAAGAAACACAGGCAAGCCCGACACATGG 5983  
Db 6033 TTCTTTTACTGTTTGTCAAAAAGGTGGAATAGAAAAAGCCTTAATGTATTGTTGAATAC 6092  
Qy 5984 GTTGAATGGGCTCTTTGAGTCAAGGCTTTTGTAGTTGTAGCACTCATCAATAGTT----- 6037  
Db 6093 ATGGTTCAAAGTCAATTTGAGTAGAGATGTTTTTAATCAGGAGTGTCCAATCATTTGGCTT 6152  
Qy 6038 -----GATCATGTGTCAGGTGGAGGCTTACCTGTGAGCCGAGCCCTG 6079  
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Qy 6167 CCCCACTTAAATTTAATTTGACAAAAGACTGTGTAATTTGTGGGATACAGTGTGATAATTG 6226  
Db 6333 AGCTCCTTATAGTAATCTGTCTATAGTAGTTTGGAGCTGCAAAACAGCCCAAGCATTA 6392  
Qy 6227 A-----TCTATGTGTGCTTTGTG 6244  
Db 6393 ATGGGTGGCACTCGGATCCCCAGATCCAGGCTCACTTCACTCTCTCTGCTCTGTTA 6452  
Qy 6245 CAAGTTCAATAGATAGATTAATAGGCCCATCAACAGCTTTTATGGGTGGAATGCAAG 6304  
Db 6453 AGAAGGGTGGTCAACTCTCTGCCAGCTTTTAAACAGCTTTTATAGTGTGAGGTGCACC 6512  
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Qy 6365 GCAAAATCATATTACTCATGCTTAAATAATCATTTATGTTGATTTAATCTTTTATAGAA 6424  
Db 6572 GCAAAATCATACATACTATAAGGATA-----TTACTATGAATGTTTACAAAT 6619  
Qy 6425 GGCTGATTAATGTTTGGTGTCTCAGAACCAATGTCAACAGCTCTTTCTAAGTGTAC 6484  
Db 6620 GCTTAAACTCTGGTTTTCTGTCTCCATCAACTAATCTTGCAATTTCT- - -AATTTGTCA 6676  
Qy 6485 CACTTTAGAAAATGCTACCTGTCTCAAAATTTGTTGTTGTTATTTTATTTTATAGTTGGA 6544  
Db 6677 CTTTGAAGAACATGGCATAAATGCTCAAAATPACTTTTGCATTTCTTATTTTCAAGCTTGA 6736  
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Db 6737 GAGAGTGGAGAGATCAAGCAATTTGGAGAACTGGAATTTGCTGTTTATGTCTCTGAGAAAT 6796  
Qy 6605 GCTTGGCTGTGACGAGAGAGAGCTAGAAAACGAAGAACTGTCTCTTCTGCTCTTCTTCAA 6664  
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Qy 6665 AAGAACAAATGAATCCCTGAATGGAATTTTTT-----ACTAAAGAAAGTGAAGAGTAAAC 6720  
Db 6857 AATAACAATTAGATGCCCCAAAGCGATTTTTTAAACAAAGAGAGATGGGAAGCCAA 6916  
Qy 6721 GTCCATCATCATTTAGAGATTTTCAATGAACCTGGCTCAGTTTGAAGAAAGAAATAGTGT 6780  
Db 6917 CTCCATCATGATGGTGGATTTCCAAATGAACCCCTGGTGTAGTTTACAAAGAAACCAATG 6976  
Qy 6781 CAA--GTGTCCATGAGACAG--AGGTAGACTTGTATAACCAAGAGATTCATTGACAAATA 6837  
Db 6977 CCACCTTTTGTATTAAAGCCAGAGAGGTAGACTTTTCTTAAGCATAGATATTTTATGATAACA 7036  
Qy 6838 TTTTATTTGTCACATGAG- - -ATACAACAGAAAATAATGTACTTTTAAAGAAATTTGTTT-- 6891  
Db 7037 TTTTCAATGTAACCTGGTGTCTATACAGAAAACAAATTTATTTTAAATTAATTTGTTCTTT 7096





Qy	3298	GTACAAGTACTTGTGGGGAGAGAAATCCACTGAGTACAAGTACTTGTGGCATGAGATC	3357
Db	1310	GAATAACATCTAGCTGTGGAAATGGCAATCATTTGAGTCTTAAGTTGTGTGAGGGAGGGGATGG	1369
Qy	3358	CACGTGAGTACAAGTACTTGTGGGGAGGGAATGGCACAGACAGAAAAGTTGAAGGGAAGG	3417
Db	1370	CATGGAGAGAAATTAAGAGAGAAAGTCGGAAATGGGAAGGCTTAA	1415
Qy	3418	AAGATCGAGAGCCCTCATGTTGTGGGGTGTGAAGAGTCACTCCCTTTTCCATGTGATGGAG	3477
Db	1416	-----GTCGGTGTGGGTTCGGAGACTGTTGCCCTGTGTGATCATGGGA	1460
Qy	3478	AGTTAAGAAAAACA-GTGTGTGAGTTTGATGTCTTCAGACACCCCAACTATGAACAT	3536
Db	1461	AGCCAAAAATCGGAGCGGTGAACATTTGATGCCGCTGAACATTTGAAACATATGAAAAAA	1520
Qy	3537	ATCCAGAGAGCGGCAGACTGTGGGAGACCTGGCAATTTAGGGAAGGCGC--GGCTTTT	3594
Db	1521	AGTTTGATGGAGTGGGCCCAGTAAAGGCCCTAGGACTTACTGAAGAGGGCTTAATTTT	1580
Qy	3595	CACACGAGAACTTTATGCTCATCTCTTGTGCTACACTCCCACTTTGATGAGGTTTCAGC	3654
Db	1581	CACATGAGATGTTTTATGTACATTTCTTGTCTTAAGCATGCAATTTCTCGAGATCAGAT	1640
Qy	3655	TCAGGTTTCGTTTCT-----ACCGTTCCTGGTACTGTGTGGAAC	3693
Db	1641	TGAGGTTTTATTCCTTACAGAAATTTGCATAAACTACTCCGCTCTTCCAAAAATGCAAC	1700
Qy	3694	TTCAAGTAGGATTCCTCCAAAGACGAGACAGCTCTCTGTGAAGGAGGACCTCGATTTCA	3753
Db	1701	CTCAGTAGGATTTCCCAAGATGAAGAGAGGTCTCTTGTGAAGGAAGTACTGGAATCTG	1760
Qy	3754	GTGTCTTAGAGAACGAATAGCTCAGAGAAATCTAGGTCAACGTGAATCTAGGTCAACAGC	3813
Db	1761	GCCTCAAGGGAATTCAGAGCTCAGGAATCTAGGTCACTGTGTGAATCTAGGTCAATG	1820
Qy	3814	GGGCAAAATGACTGAACGCCTCTATTTCAGGTGAACGGTCACTGGCTCAGATATACTG	3873
Db	1821	TGGCAAAATTTACTTAAGAGCTTTAAATTCAGGTGAATTTGTACTGTAACCTCCATGGGTG	1880
Qy	3874	AGTATTGGGCTCCACCGGATAGATTCGTGTAGTGA-GTCTGCTTTTATTTTTCAGCA	3932
Db	1881	GAGGTTTCATTAAGTTTCAGCACAACTAAGATAGTTATGCTGTTATTTGTTTATAGCA	1940
Qy	3933	CATCAGCGGTGAACACAGAACATCCAGAGAAGTGTCAAGGCTGAAGGAGACAGTGAA	3992
Db	1941	TATTGAAGGTGATGACCTGTCATATCCAGAGGAATGTCAAAAGCTGAAGACACAGTGAA	2000
Qy	3993	AAAGGTACTATTGGCAAGCCACAATACTAAGCCATTCAGTAG--GAGACGTGGGATTC	4050
Db	2001	AAAGGTAGGACTGATAACTGTCAATGCTAAGTCAATGCAATAGGAGACAAAATGTTGTTT	2060
Qy	4051	TTTCTCTGCTTCCAGTCCCTTCTACTTTTGTATACATTTTATTTTGAATGTGTCTACTATCTG	4110
Db	2061	TTCTTTCTCTTCTTCTTCCCATCACCTTTGTGATTTTTTCACTTGATTTCTCTCAACCAG	2120
Qy	4111	GTCCATTACTCGCTTAGCTGCACCTGATCTTAGCTGGGTCTATAGATCTTTTCAATCTGTG	4170
Db	2121	GGCAATTA----CTTTGGTGTCTGTATGTAGATATATCTATATATCTAGATGTCAGTT	2176
Qy	4171	TCTAAATTT---GTAAGTCAAAATTCGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTC	4227
Db	2177	TCCAAATCTTGCAAATTTGTAGAAATTCAGAACTGGTTGGGATCTTAGCTTGTCTAGTCA	2236
Qy	4228	ATGAGCACTTGCTCGGAGGATGGCTTGTGACAGAGTCAATGCTTAGAAGACAGCATCCCTG	4287
Db	2237	ATAACTCAGATTCCTGGGAGTGGTFCAGTGGCAGAGATAGGGCTAGAAATGCAGGTCTCCTG	2296
Qy	4288	ATTCACAGCTCTGCAC--TTGCCTAGTCGCCATGTGTAATTTACTTTTGCTTGATTAAGTAT	4346
Db	2297	AATCCCAAGCCAGCACATTTTCCCGGTGGTGATACAGATTAGTTAGTTTGTGACATTAATCT	2356
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[illegible]

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Qy 5448 CAGAGTCCCCGAGACCA-----CCACAGGTGTTAAGTAGG 5485  
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Qy 5963 AGGCAAGCCGACCAATGGGTTGAATGTTGGTCTTTGAGTCAAGGCTTTTGGTTGAG 6022  
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Qy 6023 CACTCATCAATAGTT-----GATCATGTGTCAGGTGGAGGC 6058  
Db 4042 GAGTGTCATTCATTTGGCTTCCCTGGACCACTTGAAAGAAATTTGTCTTGGTACACAT 4101  
Qy 6059 TACCTGTGAGCCGAGCCCTGCTGGCTTCGCACTTAACATCTCCAGGTCTCAGTATCACT 6118  
Db 4102 AABATACAGAACAAATAGCTGATGAGCTTAAAGAGTCCATGCATAAATCTCATACTGTTT 4161  
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Qy 6164 -----AACCCCACTTAAATTTTAAATTTGACAAAGACTGTGTAATTTG 6205  
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Qy 6206 TGGGATACAGTGTGATAATTGA----- 6227  
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RESULT 9

US-09-354-243B-25  
; Sequence 25, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa  
; TITLE OF INVENTION: (T1P8)  
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.1  
; CURRENT APPLICATION NUMBER: US/09/354,243B  
; CURRENT FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 25  
; LENGTH: 4797  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-09-354-243B-25

Query Match 9.2%; Score 686; DB 3; Length 4797;  
Best Local Similarity 53.8%; Pred. No. 3.6e-163;  
Matches 2644; Conservative 0; Mismatches 1875; Indels 393; Gaps 44;

Qy 2034 CTCTCTCTCACTTATCAACTTTGACACTTTGCGGATCTCTGATGGCTCTCTCCCTGCAGAA 2093  
Db 29 CTCCTCCCAAGTCACAGTTGCTCGAGTTAGAAATGCTCGCAATGGCCGCCCTGCAGAA 88  
Qy 2094 ATCTATCAGTTTTTCCCTTATGGGACTTTTGGCCGCGCAGCTGCTGTCTCTCATTTGCCCT 2153  
Db 89 ATCTGAGCTCTTTCCTTATGGGACCTTGGCCACACAGTGCCTCTCTCTCTTGGCCCT 148  
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Qy 2274 TCTCTTCTCTCCCATCGCTTGGCCATTTTCTCTGAAGCACTTTCGCAAACTCTTTAGGGG 2333  
Db 269 AATCTGCTCTTCTCGTTGGATCTACTTGAATCCAAATAGTTCTTAAACTTTCTTCA 328  
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Db 329 GAGCATCTCTAAGAGCTTTAGGAACCCACTGTTTATCCCTGAGGGTAGATAAAATTTCTG 388







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Db 990 ATATGGATTTATTATAGAAACATTATCTGCTATTGATATTTAGTATAAGSCAAATAATA 1049
Qy 7075 TTTATGACAATACTATGGAAAACAAGATATCTTAGGCTTTAAATAAACACATGGATATCAT 7134
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Qy 7135 AAA 7137
Db 1110 AAA 1112

RESULT 12
US-09-354-243B-7
; Sequence 7, Application US/09354243B
; Patent No. 6359117
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renauld, Jamila
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Pa
; TITLE OF INVENTION: (T1P8)
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.1
; CURRENT APPLICATION NUMBER: US/09/354,243B
; CURRENT FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 7
; LENGTH: 1119
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-09-354-243B-7

Query Match 8.1%; Score 601.4; DB 3; Length 1119;
Best Local Similarity 99.8%; Pred. No. 4.9e-142;
Matches 602; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Db 570 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGCTGCTCTTCCT 629
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Db 630 GCCTTCTTAAAGAAACAATAAGATCCCTGAATGACCTTTTACTTAAAGGAAAGTGAGAA 689
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Db 1050 TTTATGACAATACTATGGAAAACAAGATATCTTAGGCTTTAAATAAACACATGGATATCAT 1109
Qy 7135 AAA 7137
Db 1110 AAA 1112

RESULT 13
US-10-084-298-3
; Sequence 3, Application US/10084298
; Patent No. 6939545
; GENERAL INFORMATION:
; APPLICANT: Jacobs, Kenneth
; APPLICANT: Pittman, Debra
; APPLICANT: Fouser, Lynette
; APPLICANT: Spaulding, Vikki
; APPLICANT: Xuan, Dejun
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory
; TITLE OF INVENTION: Disorders
; FILE REFERENCE: GI5358 CIP
; CURRENT APPLICATION NUMBER: US/10/084,298
; CURRENT FILING DATE: 2002-09-10
; PRIOR APPLICATION NUMBER: 60/270,823
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/281,353
; PRIOR FILING DATE: 2001-04-03
; PRIOR APPLICATION NUMBER: 60/131,473
; PRIOR FILING DATE: 1999-04-28
; PRIOR APPLICATION NUMBER: 09/561,811
; PRIOR FILING DATE: 2000-04-28
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Murine
US-10-084-298-3

Query Match 8.0%; Score 598.2; DB 3; Length 1166;
Best Local Similarity 99.5%; Pred. No. 3.3e-141;
Matches 600; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 6535 ATAGCTTTGGAGAGAGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 6594
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Qy 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGCTGCTCTTCCT 6654
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Qy 6835 ATATTTTATTGCTCACTGATGATACAACAGAAAAATAATGTACTTTTAAAAAAATGTTTGA 6894
Db 833 ATATTTTATTGCTCACTGATGATACAACAGAAAAATAATGTACTTTTAAAAAAATGTTTGA 892
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RESULT 14  
US-09-178-973B-9  
; Sequence 9, Application US/09178973B  
; Patent No. 6274710  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543  
; CURRENT APPLICATION NUMBER: US/09/178,973B  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 17  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
US-09-178-973B-9  
Query Match 7.5%; Score 555.2; DB 3; Length 1111;  
Best Local Similarity 96.0%; Pred. No. 2.6e-130;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
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Db 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGACTAGAAAACGAAGAACTGCTCCTTCT 627  
QY 6655 GCCTTCTAAAAGAACCAATTAAGATCCCTGAATGGAATTTTATTAAGGAAAGTGAGAA 6714  
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Db 1108 TAAA 1111  
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; Sequence 9, Application US/09419568P  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/419,568P  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-419-568P-9  
Query Match 7.5%; Score 555.2; DB 3; Length 1111;  
Best Local Similarity 96.0%; Pred. No. 2.6e-130;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
QY 6535 ATAGCTTGGAGAGAGTGGAGAGATCAAGGCGATTGGGGAACCTGACCTGCTGTTTATGTC 6594  
Db 508 AAGCTTGGAGAGAGCGGAGAGATCAAGCGATCGGGGAACCTGACCTGCTGTTTATGTC 567  
QY 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGACTAGAAAACGAAGAACTGCTCCTTCT 6654  
Db 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGACTAGAAAACGAAGAACTGCTCCTTCT 627  
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QY 6835 ATATTTTATTTGTCATGATGATCAACAGAAAAAATATGTAATTTAAAAAATTTGTTGAA 6894  
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Qy	7015	ATATGGATTTTATTATAGAAACATTTCTGCTATTGATATTT-AGTATAAGGCAAAATAAT	7073
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Qy	7134	TAAA	7137
Db	1108	TAAA	1111

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Job time : 932.71 secs



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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:16 ; Search time 3723.38 Seconds  
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16534.837 Million cell updates/sec

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Perfect score: 7445  
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Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 9793542 seqs, 4134689005 residues

Total number of hits satisfying chosen parameters: 19587084

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Published Applications NA.Main.\*  
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	7445	100.0	7445	7	US-10-627-273-8
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4	4245.2	57.0	5935	7	US-10-627-273-29
5	686	9.2	4797	3	US-09-751-797-25
6	686	9.2	4797	7	US-10-627-273-25
7	601.4	8.1	1119	7	US-09-751-797-7
8	601.4	8.1	1119	3	US-10-627-273-7
9	598.2	8.0	1166	5	US-10-084-298-3
10	598.2	8.0	1166	6	US-10-256-977-3
11	598.2	8.0	1166	8	US-10-873-972-3
12	598.2	8.0	1166	10	US-11-157-387-3
13	555.2	7.5	1111	3	US-09-751-797-9
14	555.2	7.5	1111	7	US-10-627-273-9
15	499	6.7	1050	5	US-10-090-365-40
16	499	6.7	1050	5	US-10-104-919-42
17	499	6.7	1050	8	US-10-807-837-10
18	499	6.7	1050	9	US-10-968-432-42
19	499	6.7	1050	10	US-11-045-944-40
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21	272.4	3.7	778	7	US-10-395-741B-37
22	272.4	3.7	778	7	US-10-806-294-37
23	217.6	2.9	1177	10	US-11-013-920-1

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25	217.6	2.9	1191	5	US-10-084-298-1	Sequence 1, Appli
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27	217.6	2.9	1191	8	US-10-873-972-1	Sequence 1, Appli
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37	215.6	2.9	1116	9	US-10-968-432-14	Sequence 14, Appli
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39	215.6	2.9	1116	10	US-11-045-944-14	Sequence 14, Appli
40	214.2	2.9	1152	3	US-09-870-574-1	Sequence 1, Appli
41	214.2	2.9	1152	5	US-10-006-867-153	Sequence 153, App
42	214.2	2.9	1152	5	US-10-066-500-125	Sequence 125, App
43	214.2	2.9	1152	5	US-10-063-547-153	Sequence 153, App
44	214.2	2.9	1152	5	US-10-063-551-153	Sequence 153, App
45	214.2	2.9	1152	5	US-10-063-616-153	Sequence 153, App

ALIGNMENTS

RESULT 1

US-09-751-797-8  
; Sequence 8, Application US/09751797  
; Patent No. US20010024652A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE REFERENCE: (TIFS) The Proteins Encoded, and Uses Thereof  
; CURRENT APPLICATION NUMBER: US/09/751,797  
; CURRENT FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-751-797-8

Query Match	100.0%;	Score 7445;	DB 3;	Length 7445;
Best Local Similarity	100.0%;	Pred. No. 0;		
Matches 7445;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
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Qy	61	GCAACACGAGCAGTATTTATACATGGTCTTCGACCATGCGAGGTACAGAGTGAATGG	120	
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Qy	121	TAAAGGCGGTATATCAGCATTAACCAACATGTTTCTTCTTCTTCTGCGCAAGCAACT	180	
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Qy	181	TGAAATCTATGCTTTAAACAATCTTCAAGCTCTTAATATAGTGTCTACAGCTGGAGTCCG	240	
Db	181	TGAAATCTATGCTTTAAACAATCTTCAAGCTCTTAATATAGTGTCTACAGCTGGAGTCCG	240	
Qy	241	CTGCTGTCCAACAGAGCTCTTGACAGCTCTCCTCTGTTTGCAATTTTATGTTCTTTGA	300	

Db	241	 CTGCTGTCACACAGAGCTCTTGAGCAGCGTCTCCTCTGTTTGCAAATTTATGTTCTTTTGA	300
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Db	301	 TCGACTCCCAACCTCTCACTTCCTGGCTCGTGATGGCCACCTTTCAACTTCTTGCATTTA	360
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Db	1021	 GGATTCAGTGTACATACAAATGCCAAAAATCCAGTATTTTGTAAATTTCTCTTCTCAACT	1080
Qy	1081	 ATCCATCTATATAGTATGTTATTCGAGCTCATTTTAAAAATAATATTTTCAGACTTTATGC	1140
Db	1081	 ATCCATCTATATAGTATGTTATTCGAGCTCATTTTAAAAATAATATTTTCAGACTTTATGC	1140
Qy	1141	 TTGCACAAGTAAAAATGTCAGAGAAATTAGCAAAATGATATTTATTTTAAAAAAA	1200
Db	1141	 TTGCACAAGTAAAAATGTCAGAGAAATTAGCAAAATGATATTTATTTTAAAAAAA	1200
Qy	1201	 TCTATGCTTTAAAAATGCTATTTAGATTTGTTCACTACCGATATTTCCAAACTTAACCTTGACC	1260
Db	1201	 TCTATGCTTTAAAAATGCTATTTAGATTTGTTCACTACCGATATTTCCAAACTTAACCTTGACC	1260
Qy	1261	 TTGGCTATGATTTCAACCTTTGATTTTGCATCTACCATTAACAGTCTCTGGAACAGAACAT	1320
Db	1261	 TTGGCTATGATTTCAACCTTTGATTTTGCATCTACCATTAACAGTCTCTGGAACAGAACAT	1320
Qy	1321	 TCTGTGGCAATGGGAGCTGTGAAGAAAGCCAACTTCTTATTTAAAAAAAACAGCTA	1380

1321	Db	TCTGTGCGCAATGGGAGCTGTGAAGAAAGCCAACTCTCTTATTAAAAAATAAAACAGCTA	1338
1381	Qy	GTTATAGTTTAGGATTCCATATACTAAAAAATAGAGATATAATTATTTTAAAAATTTGA	1440
1381	Db		
1381	Db	GTTATAGTTTAGGATTCCATATACTAAAAAATAGAGATATAATTATTTTAAAAATTTGA	1440
1441	Qy	AATAAATCTCAAAGTTTTCATTATGGCTTATTTCAAAGCACAGAAATATAGACACGGGTCT	1500
1441	Db		
1441	Db	AATAAATCTCAAAGTTTTCATTATGGCTTATTTCAAAGCACAGAAATATAGACACGGGTCT	1500
1501	Qy	TTTATTTCTCGTCACCTCTAAAGAGATAAGAAATCTATGAAGTTTGGTGGGAAAAATGAGTCC	1560
1501	Db		
1501	Db	TTTATTTCTCGTCACCTCTAAAGAGATAAGAAATCTATGAAGTTTGGTGGGAAAAATGAGTCC	1560
1561	Qy	GTGACCAAAACGGTGACTCAATAGCTACGGGAGATCAAAAGCGTGCTCTACTCAATACAGAA	1620
1561	Db		
1561	Db	GTGACCAAAACGGTGACTCAATAGCTACGGGAGATCAAAAGCGTGCTCTACTCAATACAGAA	1620
1621	Qy	TCTACTACGGCAAGCCATGGCTTCTTTTGAAACCGTGTTTAAAGATTTCTGGGATTTT	1680
1621	Db		
1621	Db	TCTACTACGGCAAGCCATGGCTTCTTTTGAAACCGTGTTTAAAGATTTCTGGGATTTT	1680
1681	Qy	GTGTGCAAAAGCACCTTGTTGGCCCTCACCGTCAGCGTTTTTAGGGAAGACTTCCCATCTCT	1740
1681	Db		
1681	Db	GTGTGCAAAAGCACCTTGTTGGCCCTCACCGTCAGCGTTTTTAGGGAAGACTTCCCATCTCT	1740
1741	Qy	CAAGGTGGGAAGGCTTGGAGGTGTGTCTTGTGGCCCTCTATGGTGGTTAGGTACTTCTC	1800
1741	Db		
1741	Db	CAAGGTGGGAAGGCTTGGAGGTGTGTCTTGTGGCCCTCTATGGTGGTTAGGTACTTCTC	1800
1801	Qy	AGAAGACAGGATGTGGAAATTAGATAATGTCTGATGTCAATATCAATACAAATACCAAAAA	1860
1801	Db		
1801	Db	AGAAGACAGGATGTGGAAATTAGATAATGTCTGATGTCAATATCAATACAAATACCAAAAA	1860
1861	Qy	ACCTGTGTGTCGGATGGCTATAAAGACGACACTTCTGCTCTCCCATCACAGACAGAG	1920
1861	Db		
1861	Db	ACCTGTGTGTCGGATGGCTATAAAGACGACACTTCTGCTCTCCCATCACAGACAGAG	1920
1921	Qy	ACACTAAACAGGTAAGCACTCAGACCTCTACAGACAATCATCTGCTTGGTACCATGCTA	1980
1921	Db		
1921	Db	ACACTAAACAGGTAAGCACTCAGACCTCTACAGACAATCATCTGCTTGGTACCATGCTA	1980
1981	Qy	CCCGACGAACAATGCTCCCTCGATGTTTTTGGCTTTTGGCTCTCTCACTAAACAGGCTCTCT	2040
1981	Db		
1981	Db	CCCGACGAACAATGCTCCCTCGATGTTTTTGGCTTTTGGCTCTCTCACTAAACAGGCTCTCT	2040
2041	Qy	CTCATTTATCAACTGTGTGACACTTGTGGGATCTCTGATGGCTGTCTGCAGAAATCTATG	2100
2041	Db		
2041	Db	CTCATTTATCAACTGTGTGACACTTGTGGGATCTCTGATGGCTGTCTGCAGAAATCTATG	2100
2101	Qy	AGTTTTTCCCTTATGGGACATTTTGGCCGCGCAGCTGCTGCTTCTCATTTGCCCTGTGGCC	2160
2101	Db		
2101	Db	AGTTTTTCCCTTATGGGACATTTTGGCCGCGCAGCTGCTGCTTCTCATTTGCCCTGTGGCC	2160
2161	Qy	CAGGAGGCAATGCGCTGCCGTCACACCCGGTGCAAACCGGTCGAAGCTTGAGGTGTCCAACTTCAG	2220
2161	Db		
2161	Db	CAGGAGGCAATGCGCTGCCGTCACACCCGGTGCAAACCGGTCGAAGCTTGAGGTGTCCAACTTCAG	2220
2221	Qy	CAGCGGTACATGCTCAACCGCACCTTTATGCTGGCCGCAAGGATACAGCTGCATCTCTTT	2280
2221	Db		
2221	Db	CAGCGGTACATGCTCAACCGCACCTTTATGCTGGCCGCAAGGATACAGCTGCATCTCTTT	2280
2281	Qy	CTCTCCATACCGCTGTGCATTTTCTGTAAGCACTTGCAAACTTTTAGGGGCGCTTTA	2340
2281	Db		
2281	Db	CTCTCCATACCGCTGTGCATTTTCTGTAAGCACTTGCAAACTTTTAGGGGCGCTTTA	2340
2341	Qy	TCTCCGCAAGTCTCACTACCTATGTTTTTGTCTCTTTTAGAGACTCTTTAAGGACTGGGT	2400
2341	Db		
2341	Db	TCTCCGCAAGTCTCACTACCTATGTTTTTGTCTCTTTTAGAGACTCTTTAAGGACTGGGT	2400
2401	Qy	CTTTTTCTATTTCTATTTCAAAGTCTCAGGACCAATTTCTTATCTTGGCTTCAGGACACA	2460
2401	Db		
2401	Db	CTTTTTCTATTTCTATTTCAAAGTCTCAGGACCAATTTCTTATCTTGGCTTCAGGACACA	2460

QY	2461	TATACCTGAATTTTATCTACAGAGCGCGCATTTAGAAAGCCACCAACGACTGCAATATCTTTC	2520
Db	2461	TATACCTGAATTTTATCTACAGAGCGCGCATTTAGAAAGCCACCAACGACTGCAATATCTTTC	2520
QY	2521	CATTTCTCTGTGCTCTCTCTGTAACCTACATCTCTTGGGTACTCTCTGAGACCCACTGCG	2580
Db	2521	CATTTCTCTGTGCTCTCTCTGTAACCTACATCTCTTGGGTACTCTCTGAGACCCACTGCG	2580
QY	2581	GACATACATCTCTACCTTACAGGCTTTTCTTCCATCTCTCTGTGTCACCCAGGACCTTAGGGT	2640
Db	2581	GACATACATCTCTACCTTACAGGCTTTTCTTCCATCTCTCTGTGTCACCCAGGACCTTAGGGT	2640
QY	2641	TTTCTCTCTTTTACGGCAGCCTTGCAGATAACACACAGACGTCCTCGGCTCATCGGGAGA	2700
Db	2641	TTTCTCTCTTTTACGGCAGCCTTGCAGATAACACACAGACGTCCTCGGCTCATCGGGAGA	2700
QY	2701	AACCTGTTCCGAGGAGTCAGTGAAGTCTCTCACTGTGATGACGAGGCTAGCTCGGGAGC	2760
Db	2701	AACCTGTTCCGAGGAGTCAGTGAAGTCTCTCACTGTGATGACGAGGCTAGCTCGGGAGC	2760
QY	2761	TGCTGACCCCTCTGGGATAGTCTGACGTATGACCCCTGCTCTTCTGTCTACCTCGCAGG	2820
Db	2761	TGCTGACCCCTCTGGGATAGTCTGACGTATGACCCCTGCTCTTCTGTCTACCTCGCAGG	2820
QY	2821	CTAAAGATCAGTCTACCTGATGAAGCAGGTGCTCAACTTTCACCTGGAAGACGTTCTGC	2880
Db	2821	CTAAAGATCAGTCTACCTGATGAAGCAGGTGCTCAACTTTCACCTGGAAGACGTTCTGC	2880
QY	2881	TCCCCAGTCAGACAGGTTCCAGCCCTACATGACGAGGTTGATACCTTCTCTGACCAAC	2940
Db	2881	TCCCCAGTCAGACAGGTTCCAGCCCTACATGACGAGGTTGATACCTTCTCTGACCAAC	2940
QY	2941	TCAGCAATCAGCTCAGCTCTGTGTAAGTCTGACTCTGCTGCTACCTATGCTCTCTCTT	3000
Db	2941	TCAGCAATCAGCTCAGCTCTGTGTAAGTCTGACTCTGCTGCTACCTATGCTCTCTCTT	3000
QY	3001	CCTCTTCTATTTCCAGTAAGAACCCGAGGTCCTGCTCTCTCTCTTTTCCAAAGAGTGAGGA	3060
Db	3001	CCTCTTCTATTTCCAGTAAGAACCCGAGGTCCTGCTCTCTCTCTTTTCCAAAGAGTGAGGA	3060
QY	3061	GGGCTCAGCACACCAACCATATAGGCCACTTGAATAGGTACAAAGGCTTTGGCTTC	3120
Db	3061	GGGCTCAGCACACCAACCATATAGGCCACTTGAATAGGTACAAAGGCTTTGGCTTC	3120
QY	3121	AATTGAGTAATATCTTTGAGTTTGTATGAGTGAAGCTTTATTTGTTTATCCATGGAAAGA	3180
Db	3121	AATTGAGTAATATCTTTGAGTTTGTATGAGTGAAGCTTTATTTGTTTATCCATGGAAAGA	3180
QY	3181	AATCAACTCAAAATCTGTAGGATGAGAAAGATGTTGGAAACGAAAAAGGCTTAGATAGA	3240
Db	3181	AATCAACTCAAAATCTGTAGGATGAGAAAGATGTTGGAAACGAAAAAGGCTTAGATAGA	3240
QY	3241	GAAACAGATCTGTGAGTATAGTACTTATGGGGGAGCAGGGGGCGATATCCACTGAGTA	3300
Db	3241	GAAACAGATCTGTGAGTATAGTACTTATGGGGGAGCAGGGGGCGATATCCACTGAGTA	3300
QY	3301	CAAGTACTTGTGGGAGAGAAATCCACTGAGTACAAGTACTCTGTGSCATGGAGATCCAC	3360
Db	3301	CAAGTACTTGTGGGAGAGAAATCCACTGAGTACAAGTACTCTGTGSCATGGAGATCCAC	3360
QY	3361	TGAGTACAAGTACTTGTGGGGGAGGGAATGGCACAGAGCAAAAGTTGAAGGGAAGGAAG	3420
Db	3361	TGAGTACAAGTACTTGTGGGGGAGGGAATGGCACAGAGCAAAAGTTGAAGGGAAGGAAG	3420
QY	3421	ATGGAGAGGCTCATGTTGGGGGTGTGAAGGTCACTCTCTTTTCCATGTGATGGAGGT	3480
Db	3421	ATGGAGAGGCTCATGTTGGGGGTGTGAAGGTCACTCTCTTTTCCATGTGATGGAGGT	3480
QY	3481	TAAAGAAACCACTGTGAGTTGTGCTTTCAGCACCCCCCAACTATGAAACATATCC	3540
Db	3481	TAAAGAAACCACTGTGAGTTGTGCTTTCAGCACCCCCCAACTATGAAACATATCC	3540

QY	3541	ACAGAGAGCGGCGAGACTGTGGGAGACCTGGGCATTTTAGGAAAGCGCGGCTTTTTCACACG	3600
Db	3541	ACAGAGAGCGGCGAGACTGTGGGAGACCTGGGCATTTTAGGAAAGCGCGGCTTTTTCACACG	3600
QY	3601	AGAAACCTTTATGCTCATCTCTTGTGTACACTCCACCTTTGATGAGTTTTCAGCTCAGGT	3660
Db	3601	AGAAACCTTTATGCTCATCTCTTGTGTACACTCCACCTTTGATGAGTTTTCAGCTCAGGT	3660
QY	3661	TTGCTTTTACCGTCTTCTGCTACTGTTGGAACCTTTCAGTAGGATTTCCCAAGACGAGGA	3720
Db	3661	TTGCTTTTACCGTCTTCTGCTACTGTTGGAACCTTTCAGTAGGATTTCCCAAGACGAGGA	3720
QY	3721	CAGCTCTTCTGTAAGGAGGACCTCGATTTTTCAGTGTCTTAGAGACGAAATAGCTCAGA	3780
Db	3721	CAGCTCTTCTGTAAGGAGGACCTCGATTTTTCAGTGTCTTAGAGACGAAATAGCTCAGA	3780
QY	3781	GAATCTAGGTCACACGTGAATCTAGGTCACAGCGGGCAAAATGACTGAACGCTCTATT	3840
Db	3781	GAATCTAGGTCACACGTGAATCTAGGTCACAGCGGGCAAAATGACTGAACGCTCTATT	3840
QY	3841	CCAGGTGAACCGGTCACTGCTCAGATATACCTGAGGTATTGGGCTCCACCGGATAAGAT	3900
Db	3841	CCAGGTGAACCGGTCACTGCTCAGATATACCTGAGGTATTGGGCTCCACCGGATAAGAT	3900
QY	3901	TCGTTAGTGAAGTCTGCTTTTATTTTTCAGACCATCAGCGGTGACGACGAAATCCAG	3960
Db	3901	TCGTTAGTGAAGTCTGCTTTTATTTTTCAGACCATCAGCGGTGACGACGAAATCCAG	3960
QY	3961	AGAAATGTCAGAGGCTGAGGAGACAGTGAAGAGTACTATTGGCAAGCCCAATACT	4020
Db	3961	AGAAATGTCAGAGGCTGAGGAGACAGTGAAGAGTACTATTGGCAAGCCCAATACT	4020
QY	4021	AAGCCATTTCAGTAGGAGAGCTGGGGATTTCTTCTCTGCTTCCAGTCCCTTCTACTTTG	4080
Db	4021	AAGCCATTTCAGTAGGAGAGCTGGGGATTTCTTCTCTGCTTCCAGTCCCTTCTACTTTG	4080
QY	4081	TAACTTTTATTTGACTGTTGCTACTATCTGCTGCTCAATCTGCTGCTGCTGCTGATC	4140
Db	4081	TAACTTTTATTTGACTGTTGCTACTATCTGCTGCTCAATCTGCTGCTGCTGCTGATC	4140
QY	4141	TAGCTGGGCTATAGATCTTCAATCTGCTTAAATTTTGAAGTCAAAATTCCTGGAGCT	4200
Db	4141	TAGCTGGGCTATAGATCTTCAATCTGCTTAAATTTTGAAGTCAAAATTCCTGGAGCT	4200
QY	4201	AGCAGAAAGCTTAGCTCAGCCAGTCTCATGAGCACTTGTCTGGAGGATGCTTGTGACAG	4260
Db	4201	AGCAGAAAGCTTAGCTCAGCCAGTCTCATGAGCACTTGTCTGGAGGATGCTTGTGACAG	4260
QY	4261	AGTCAATGCTAGAAGACAGCATCCCTGATTTCCAGCTCTGCACTTGTGCTAGTGGCCATGT	4320
Db	4261	AGTCAATGCTAGAAGACAGCATCCCTGATTTCCAGCTCTGCACTTGTGCTAGTGGCCATGT	4320
QY	4321	GTAAATTTTGGCTTGTATGATTTTGGAAAGCCAGTTCCCGAGGCTCATCAATATC	4380
Db	4321	GTAAATTTTGGCTTGTATGATTTTGGAAAGCCAGTTCCCGAGGCTCATCAATATC	4380
QY	4381	TGAAGAACCATGCTATTTGAAAACTAGAAAAGCTGGGCAAAACTTACTAGAGATGATTTTG	4440
Db	4381	TGAAGAACCATGCTATTTGAAAACTAGAAAAGCTGGGCAAAACTTACTAGAGATGATTTTG	4440
QY	4441	AGCTCAATTAACCGGATGCTCTGAAATGTGGCAAAATCAACCCAGATAACAAAGAG	4500
Db	4441	AGCTCAATTAACCGGATGCTCTGAAATGTGGCAAAATCAACCCAGATAACAAAGAG	4500
QY	4501	CTGGATTTTGAATAGGACAGTATTTAGATCACTGGTATTTAATAGCTATCATCTTAAT	4560
Db	4501	CTGGATTTTGAATAGGACAGTATTTAGATCACTGGTATTTAATAGCTATCATCTTAAT	4560
QY	4561	TAAATATAGGGCTATATATATTTAAGATTAAACACAAAGAGTGGATAGCTTCCCAAT	4620
Db	4561	TAAATATAGGGCTATATATATTTAAGATTAAACACAAAGAGTGGATAGCTTCCCAAT	4620
QY	4621	TTACTTTGGCTGTTTCAAAAGAGTAAAAATATCAGTCACTGGATTAAATATAGTGTCTATG	4680

4621	Db	 TTACTTGGCCTGGT	 TTCAAAGAGTAAAA	 TATCAGTCATG	 TAAATATATAGT	 TATAGTGCATG	4680
4681	Qy	 AAAGTAGAGATG	 GAAGAACCCCTT	 CTTCTACCTTTT	 TACCTTCAATTT	 CTTCTAGT	4740
4681	Db	 AAAGTAGAGATG	 GAAGAACCCCTT	 CTTCTACCTTTT	 TACCTTCAATTT	 CTTCTAGT	4740
4741	Qy	 TTCACACCCCTG	 ATCAAGCCACTAG	 TAGTAAGCACCT	 ATCTGCTGAGCT	 ATTATATAGCTTT	4800
4741	Db	 TTCACACCCCTG	 ATCAAGCCACTAG	 TAGTAAGCACCT	 ATCTGCTGAGCT	 ATTATATAGCTTT	4800
4801	Qy	 ACAGCAAAACAA	 CATTTGCTGTG	 GCGCTTTT	 TTGGGAAGGAA	 CAGAGATCAGGAGGCTC	4860
4801	Db	 ACAGCAAAACAA	 CATTTGCTGTG	 GCGCTTTT	 TTGGGAAGGAA	 CAGAGATCAGGAGGCTC	4860
4861	Qy	 AGGCTAGCAAGT	 CTGACTCGCCCT	 TAAGCCAGAGG	 CATGCTCATAGC	 AGAGAGTGAG	4920
4861	Db	 AGGCTAGCAAGT	 CTGACTCGCCCT	 TAAGCCAGAGG	 CATGCTCATAGC	 AGAGAGTGAG	4920
4921	Qy	 GCTCTTCGCAAG	 TGGGTGCTTAA	 GTAATCAGAAA	 CAGGAAGCCT	 CCGGTTCGATGGAAT	4980
4921	Db	 GCTCTTCGCAAG	 TGGGTGCTTAA	 GTAATCAGAAA	 CAGGAAGCCT	 CCGGTTCGATGGAAT	4980
4981	Qy	 TATCAGTAAGAT	 ATCTACCCCTTA	 TCTCCTTCTA	 TCGAACTTAAT	 CGTCTCTTTTCTTG	5040
4981	Db	 TATCAGTAAGAT	 ATCTACCCCTTA	 TCTCCTTCTA	 TCGAACTTAAT	 CGTCTCTTTTCTTG	5040
5041	Qy	 TGCTAGGCTGAT	 ATAACACTTGTT	 CTTTTGGAGT	 GTTTCATGGCTT	 TGATGATTTTA	5100
5041	Db	 TGCTAGGCTGAT	 ATAACACTTGTT	 CTTTTGGAGT	 GTTTCATGGCTT	 TGATGATTTTA	5100
5101	Qy	 GTGCTCTGCCAG	 TTCCTTTAGAGG	 TGTGTACTCT	 TGACCTGGCT	 TGGATGTTAGC	5160
5101	Db	 GTGCTCTGCCAG	 TTCCTTTAGAGG	 TGTGTACTCT	 TGACCTGGCT	 TGGATGTTAGC	5160
5161	Qy	 ATGCCAAAGGCA	 CACTTCTGATG	 CCTGTGTAAA	 AGGTTATTTAT	 CTATTACTTTGTC	5220
5161	Db	 ATGCCAAAGGCA	 CACTTCTGATG	 CCTGTGTAAA	 AGGTTATTTAT	 CTATTACTTTGTC	5220
5221	Qy	 TTTGGAAAGGTG	 AAGCGTGTGAGA	 AAGAACTCAG	 CAGGAGATGT	 CTCTGTAGAAA	5280
5221	Db	 TTTGGAAAGGTG	 AAGCGTGTGAGA	 AAGAACTCAG	 CAGGAGATGT	 CTCTGTAGAAA	5280
5281	Qy	 ACTTTTTTTTTT	 CCCCCTTAAAT	 GCGCTTAAT	 CCACTTCAGT	 CAACTTGTGATTTAT	5340
5281	Db	 ACTTTTTTTTTT	 CCCCCTTAAAT	 GCGCTTAAT	 CCACTTCAGT	 CAACTTGTGATTTAT	5340
5341	Qy	 ATGCTGTCAAT	 GAAAGAGTGT	 TTAGGCCGCT	 CTCATGGCT	 CTGGGAAAGCACCAATA	5400
5341	Db	 ATGCTGTCAAT	 GAAAGAGTGT	 TTAGGCCGCT	 CTCATGGCT	 CTGGGAAAGCACCAATA	5400
5401	Qy	 GGGGAAGGAAT	 GTGTATGCTGAGA	 AATCTGAC	 CGGCAGGAAA	 CTGGTCTCAGAGCTCCCCG	5460
5401	Db	 GGGGAAGGAAT	 GTGTATGCTGAGA	 AATCTGAC	 CGGCAGGAAA	 CTGGTCTCAGAGCTCCCCG	5460
5461	Qy	 AAGACCAACC	 CAGCTGTAA	 GTAGTGAAG	 CAAGTCTCAGG	 GTGGCTCATCTAATAGATG	5520
5461	Db	 AAGACCAACC	 CAGCTGTAA	 GTAGTGAAG	 CAAGTCTCAGG	 GTGGCTCATCTAATAGATG	5520
5521	Qy	 CAGAGCGAGG	 GAAGATAGCT	 TACAAAGTTT	 CATAGGGT	 CCGGAGCTTTAAAGATAC	5580
5521	Db	 CAGAGCGAGG	 GAAGATAGCT	 TACAAAGTTT	 CATAGGGT	 CCGGAGCTTTAAAGATAC	5580
5581	Qy	 TAGCTGCTTGG	 GCTTATAAC	 AAGAGTCTG	 GGAAGGCAAG	 GTGAGGGAATGG	5640
5581	Db	 TAGCTGCTTGG	 GCTTATAAC	 AAGAGTCTG	 GGAAGGCAAG	 GTGAGGGAATGG	5640
5641	Qy	 AAAGGGAAGG	 AAAAAGATGT	 ATAGAGCA	 CTTGGAACAG	 CTCTACACAGATTT	5700
5641	Db	 AAAGGGAAGG	 AAAAAGATGT	 ATAGAGCA	 CTTGGAACAG	 CTCTACACAGATTT	5700
5701	Qy	 TTCCTTGGA	 CAATCTAGAA	 GGTAGTGGAT	 TAGGTCATTG	 CAGGGGAGCTTGTCTTGG	5760

5701	Db	TTCTTGGAAACAATCTATGAAAGGTAGTGGATTTAGGTGATTGACGGGGGACCTTGTCTTTTGGCCAT	5761
5761	Qy	TTGAAATCTGGGTTTTTGTCTCTCCATTTAGAGTTTGAAGGGTCAACCCCTTTTATACCCTCGAA	5820
5761	Db	TTGAAATCTGGGTTTTTGTCTCTCCATTTAGAGTTTGAAGGGTCAACCCCTTTTATACCCTCGAA	5820
5821	Qy	TGAGGAGGAAGAAGGGGTGTATGACTCTTACCTGGAGTTTACTAGTTTACGCAATG	5880
5821	Db	TGAGGAGGAAGAAGGGGTGTATGACTCTTACCTGGAGTTTACTAGTTTACGCAATG	5880
5881	Qy	GAACAGACACTCGGGACCTCCTCTTGACAAAAAATAATGGAAACCTGTGTCTCTTGT	5940
5881	Db	GAACAGACACTCGGGACCTCCTCTTGACAAAAAATAATGGAAACCTGTGTCTCTTGT	5940
5941	Qy	TGTTCTTTTGTTAAGAAAGCACAGGCAAAAGCCCGAACCATATGGGTTTGAATGTGGGTCCTT	6000
5941	Db	TGTTCTTTTGTTAAGAAAGCACAGGCAAAAGCCCGAACCATATGGGTTTGAATGTGGGTCCTT	6000
6001	Qy	GAGTCAAGGCTTTTGAGTTGAGCACTCATCAATPAGTTTGATCATGGTCAAGTGGAGGGCTA	6060
6001	Db	GAGTCAAGGCTTTTGAGTTGAGCACTCATCAATPAGTTTGATCATGGTCAAGTGGAGGGCTA	6060
6061	Qy	CCTGTCAAGCCGAGCCCTGCTGGCTTCGCACATTAACATCTCCAGGTCTCAGTATCACTTC	6120
6061	Db	CCTGTCAAGCCGAGCCCTGCTGGCTTCGCACATTAACATCTCCAGGTCTCAGTATCACTTC	6120
6121	Qy	CTCCTACTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTTTCCAAACCCCACTAAATTT	6180
6121	Db	CTCCTACTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTTTCCAAACCCCACTAAATTT	6180
6181	Qy	AATTGCAAAAGACTGTGTAAATTTGTGGGATACAGTGTGATAATGATCTATGTGTGCAT	6240
6181	Db	AATTGCAAAAGACTGTGTAAATTTGTGGGATACAGTGTGATAATGATCTATGTGTGCAT	6240
6241	Qy	TGTGCAAGGTTCAATAAGATAGATTAATAGGCCCATCAACAGCTTTATGGGTGTGAATG	6300
6241	Db	TGTGCAAGGTTCAATAAGATAGATTAATAGGCCCATCAACAGCTTTATGGGTGTGAATG	6300
6301	Qy	CAAGTAATPAGGTAGATGCTGTGGTGTCTTTAGTTCAGAAAGGCATGATTTTAAGGTC	6360
6301	Db	CAAGTAATPAGGTAGATGCTGTGGTGTCTTTAGTTCAGAAAGGCATGATTTTAAGGTC	6360
6361	Qy	TTGGGCAAAATCATATTACTCATGCTTAAATAATACATTAATGTTGATTTAAATCTTTTAG	6420
6361	Db	TTGGGCAAAATCATATTACTCATGCTTAAATAATACATTAATGTTGATTTAAATCTTTTAG	6420
6421	Qy	AGAAGGCTGATCTATGGTTTTGTGTCTCAGCAAGCAAAATGTCACCAGCTCTTTCTAACTG	6480
6421	Db	AGAAGGCTGATCTATGGTTTTGTGTCTCAGCAAGCAAAATGTCACCAGCTCTTTCTAACTG	6480
6481	Qy	GTACCACTTTTGAATAATGCTACCTGTCTCAAAATTTGGTTTGTATCTCTTATTTTCATAGCT	6540
6481	Db	GTACCACTTTTGAATAATGCTACCTGTCTCAAAATTTGGTTTGTATCTCTTATTTTCATAGCT	6540
6541	Qy	TGGAGAGGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTCCTTGAG	6600
6541	Db	TGGAGAGGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTCCTTGAG	6600
6601	Qy	AAATGCTTGGCTGTAGCGAGAGAAAGACTAGAAAAAAGAAAGTCTCTTCCGCTTC	6660
6601	Db	AAATGCTTGGCTGTAGCGAGAGAAAGACTAGAAAAAAGAAAGTCTCTTCCGCTTC	6660
6661	Qy	TAAAAAGAAACAATAAGATCCCTGAATGGGACTTTTTTATTAAGAGAAAGTGGAGAGCTTAAC	6720
6661	Db	TAAAAAGAAACAATAAGATCCCTGAATGGGACTTTTTTATTAAGAGAAAGTGGAGAGCTTAAC	6720
6721	Qy	GTCCATCATCATTTAGAGATTTTACATATGAACCTGGCTCAGTTTGAATAAGAAAAATAGTGT	6780
6721	Db	GTCCATCATCATTTAGAGATTTTACATATGAACCTGGCTCAGTTTGAATAAGAAAAATAGTGT	6780
6781	Qy	CAAGTTGTCCATGAGACCGAGGTAGACTTGTATTAACCAAAAGATTCATTTGACAAATATTT	6840
6781	Db	CAAGTTGTCCATGAGACCGAGGTAGACTTGTATTAACCAAAAGATTCATTTGACAAATATTT	6840



Db	1021	GGATTCAAGTACATCAATGCAAAAATCCAGTATTTTGTAAATTTCTCTCTCAACT	1080
Qy	1081	ATCCATCTATATAGTATGTTATGTAGGCTCATTTAAAAATATATTTTGAGACTTATGC	1140
Db	1081	ATCCATCTATATAGTATGTTATGTAGGCTCATTTAAAAATATATTTTGAGACTTATGC	1140
Qy	1141	TTGCACAAGTAAATGTCAGAGAAATTAGCAAAATGATATAGTATATTTTAAAAAAA	1200
Db	1141	TTGCACAAGTAAATGTCAGAGAAATTAGCAAAATGATATAGTATATTTTAAAAAAA	1200
Qy	1201	TCTATGCTTAAAAATGTTCTATTTAGATGTTTCACTACCGATATTTCCAACTTTAAGC	1260
Db	1201	TCTATGCTTAAAAATGTTCTATTTAGATGTTTCACTACCGATATTTCCAACTTTAAGC	1260
Qy	1261	TTGGCTATGATTTCAACCTTTGATATTTGATCTACCAATACAGTCTCTGAACCAAGAT	1320
Db	1261	TTGGCTATGATTTCAACCTTTGATATTTGATCTACCAATACAGTCTCTGAACCAAGAT	1320
Qy	1321	TCTGTGGCAATGGGAGCTGTGAAGAAAGCAACATTTCTTATTTAAAAAAAACAGCTA	1380
Db	1321	TCTGTGGCAATGGGAGCTGTGAAGAAAGCAACATTTCTTATTTAAAAAAAACAGCTA	1380
Qy	1381	GTTATAGTTTGGATTTCCATATCTAAAAAAAATPAGAGATATATTTTAAAAAATTGA	1440
Db	1381	GTTATAGTTTGGATTTCCATATCTAAAAAAAATPAGAGATATATTTTAAAAAATTGA	1440
Qy	1441	AATTAATCTCAAGTTTCAATATGGCTTATTTTCAAGCAGAGATATAGGACCGGTCT	1500
Db	1441	AATTAATCTCAAGTTTCAATATGGCTTATTTTCAAGCAGAGATATAGGACCGGTCT	1500
Qy	1501	TTTATTTCTGGTCACTTCTAAAGAGATAAGAACTATGAAGTTGGTGGGAAATGAGTCC	1560
Db	1501	TTTATTTCTGGTCACTTCTAAAGAGATAAGAACTATGAAGTTGGTGGGAAATGAGTCC	1560
Qy	1561	GTGACCAAAACGCTGACTCAATAGCTACGGGAGATCAAAGGCTGCTCTACTCAATCAGAA	1620
Db	1561	GTGACCAAAACGCTGACTCAATAGCTACGGGAGATCAAAGGCTGCTCTACTCAATCAGAA	1620
Qy	1621	TCTACTACGGCAAGCCATGCTTTCTTTGAAAACCGTGTTTTGAAGATTTCTGGGATTT	1680
Db	1621	TCTACTACGGCAAGCCATGCTTTCTTTGAAAACCGTGTTTTGAAGATTTCTGGGATTT	1680
Qy	1681	GTGTGCAAAAGCACTTGTGGCCCTCACCGTGAGCTTTTAGGCAAGACTTCCCATCTCT	1740
Db	1681	GTGTGCAAAAGCACTTGTGGCCCTCACCGTGAGCTTTTAGGCAAGACTTCCCATCTCT	1740
Qy	1741	CAAGGTGGGAAGGCTTGGAGTGGTGTCTTGTGGCCCTCCTATGGTGGTTAGGTACTTCTC	1800
Db	1741	CAAGGTGGGAAGGCTTGGAGTGGTGTCTTGTGGCCCTCCTATGGTGGTTAGGTACTTCTC	1800
Qy	1801	AGAAAGACGGACTGGAAATTAGATAATGTCTGATGTCTATCATTTCAATACCAAAAAA	1860
Db	1801	AGAAAGACGGACTGGAAATTAGATAATGTCTGATGTCTATCATTTCAATACCAAAAAA	1860
Qy	1861	ACCTGTGTCCCGATGGCTATAAAGCAGCAACTTCTGCCCTCCCATCAAGCAGAG	1920
Db	1861	ACCTGTGTCCCGATGGCTATAAAGCAGCAACTTCTGCCCTCCCATCAAGCAGAG	1920
Qy	1921	ACACCTAAACAGGTAAGCACTCAGACCTCTACAGCAATCATCTGCTTGGTACCATGCTA	1980
Db	1921	ACACCTAAACAGGTAAGCACTCAGACCTCTACAGCAATCATCTGCTTGGTACCATGCTA	1980
Qy	2041	CCGACGAACATGCTCCCTCATGTTTGGCTTTTGTCTCTCTCACTAAAGGCTCTCCT	2100
Db	2041	CCGACGAACATGCTCCCTCATGTTTGGCTTTTGTCTCTCTCACTAAAGGCTCTCCT	2100
Qy	2101	AGTTTTCCTTATGGGACTTTGGCCGCAAGCTGCCTGCTTCTCAITGGCCCTGTGGCC	2160
Db	2101	AGTTTTCCTTATGGGACTTTGGCCGCAAGCTGCCTGCTTCTCAITGGCCCTGTGGCC	2160

Db	2101	AGTTTTCCTTATGGGACTTTGGCCGCAAGCTGCCTGCTTCTCAITGGCCCTGTGGCC	2160
Qy	2161	CAGGAGGCAAAATGCGCTGCCCGTCAACACCCGGTGCAAGCTTGAGGTGTCCAATCTCCAG	2220
Db	2161	CAGGAGGCAAAATGCGCTGCCCGTCAACACCCGGTGCAAGCTTGAGGTGTCCAATCTCCAG	2220
Qy	2221	CAGCCGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGGAGGTACAGCTGCATCTCTTT	2280
Db	2221	CAGCCGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGGAGGTACAGCTGCATCTCTTT	2280
Qy	2281	CTCTCCATACCGCTTGCCCATTTTCTCTGAAGCACTTGCAACTCTTTAGGGGGCTTTTA	2340
Db	2281	CTCTCCATACCGCTTGCCCATTTTCTCTGAAGCACTTGCAACTCTTTAGGGGGCTTTTA	2340
Qy	2341	TCTCCGCAAGTCTCACTACCTATGTTTCTGCTCTCTTTAGAGACTCTTTAAGGACTGGGT	2400
Db	2341	TCTCCGCAAGTCTCACTACCTATGTTTCTGCTCTCTTTAGAGACTCTTTAAGGACTGGGT	2400
Qy	2401	CTTTTCTATTTCTATTTCAAGGTCTCAGGACCATTTCTTATCTTGGCCTTTCAGGACACA	2460
Db	2401	CTTTTCTATTTCTATTTCAAGGTCTCAGGACCATTTCTTATCTTGGCCTTTCAGGACACA	2460
Qy	2461	TATACTGAATTTTATCTACAGAGCGCATTTAGAAAGCCACCCAGACTGCAATACTTTC	2520
Db	2461	TATACTGAATTTTATCTACAGAGCGCATTTAGAAAGCCACCCAGACTGCAATACTTTC	2520
Qy	2521	CATTTCTCTGTGCTCTCTCTGAACATCATACTCTCTTGGCTACTCTCTGAGACCCACTGCG	2580
Db	2521	CATTTCTCTGTGCTCTCTCTGAACATCATACTCTCTTGGCTACTCTCTGAGACCCACTGCG	2580
Qy	2581	GACATACATCTCTACTTTACAGGCTTTTCTTCCATCTCTTGTCAACCCAGGACCTTAGGGT	2640
Db	2581	GACATACATCTCTACTTTACAGGCTTTTCTTCCATCTCTTGTCAACCCAGGACCTTAGGGT	2640
Qy	2641	TTTCTCTCTTTTACAGGCGCTTTCAGAGATTAACACACAGCTCCGGCTCATCGGGGAGA	2700
Db	2641	TTTCTCTCTTTTACAGGCGCTTTCAGAGATTAACACACAGCTCCGGCTCATCGGGGAGA	2700
Qy	2701	AATGTTTCCGAGAGTCAGTGAAGTCTCACCTGTGATGAGCAGGCTAGCTGGGGAGC	2760
Db	2701	AATGTTTCCGAGAGTCAGTGAAGTCTCACCTGTGATGAGCAGGCTAGCTGGGGAGC	2760
Qy	2761	TGGTGAACCTCTCTGGGATAGTCTGACGTATGACCCCTGCTGCTTCTTGTCTACCTGAGG	2820
Db	2761	TGGTGAACCTCTCTGGGATAGTCTGACGTATGACCCCTGCTGCTTCTTGTCTACCTGAGG	2820
Qy	2821	CTAAAGATCAGTCTCCTGATGAGCAGTGTCTCAACTTCAACCTGGAAGAGCTTCTGC	2880
Db	2821	CTAAAGATCAGTCTCCTGATGAGCAGTGTCTCAACTTCAACCTGGAAGAGCTTCTGC	2880
Qy	2881	TCCCCCAGTCAGACAGCTTCCAGCCCTACATGACGAGGTGGTACCTTCTCTGACCAAAAC	2940
Db	2881	TCCCCCAGTCAGACAGCTTCCAGCCCTACATGACGAGGTGGTACCTTCTCTGACCAAAAC	2940
Qy	2941	TCAGCAATCAGCTCAGCTCCTGTGTAAAGTCTGACTCTGGCTACCTATGCTCTCTCTT	3000
Db	2941	TCAGCAATCAGCTCAGCTCCTGTGTAAAGTCTGACTCTGGCTACCTATGCTCTCTCTT	3000
Qy	3001	CCTCTTCTATTTCCAGTAAAGAACCCGAGGTCTCTCTCTTTCACAGAGTGAAGA	3060
Db	3001	CCTCTTCTATTTCCAGTAAAGAACCCGAGGTCTCTCTCTTTCACAGAGTGAAGA	3060
Qy	3061	GGGCTCAGCACCAACCATCATAGGCCATTTGAAATAGTTCACAAAGGCTTTGGGCTTC	3120
Db	3061	GGGCTCAGCACCAACCATCATAGGCCATTTGAAATAGTTCACAAAGGCTTTGGGCTTC	3120
Qy	3121	AATTGAGTAATCTTCTGAGTTTGTATGAGTGAAGCTTTATTTGTTTTATCCATGGAAGA	3180
Db	3121	AATTGAGTAATCTTCTGAGTTTGTATGAGTGAAGCTTTATTTGTTTTATCCATGGAAGA	3180
Qy	3181	AATCAACTCAAAATCTCTAGGATGAGAAAGATGTTGGGAACGAAAGGCTCTAGATAGA	3240
Db	3181	AATCAACTCAAAATCTCTAGGATGAGAAAGATGTTGGGAACGAAAGGCTCTAGATAGA	3240



Qy	3241	GAACAGATCTGCTGAGTACTATGATGGGGGAGCAGGGGGCGATATCCACTGAGTA	3300
Dd	3241	GAACAGATCTGCTGAGTACTATGATGGGGGAGCAGGGGGCGATATCCACTGAGTA	3300
Qy	3301	CAAGTACTTGTGGGGAGAGAAATCCACTGAGTACAAGTACTTGTGTGGCATGGAGATCCAC	3360
Dd	3301	CAAGTACTTGTGGGGAGAGAAATCCACTGAGTACAAGTACTTGTGTGGCATGGAGATCCAC	3360
Qy	3361	TGAGTACAAGTACTTGTGGGGGAGGGAATGGCACAGAGCAAAAGTTGAAGGGAAGGAAG	3420
Dd	3361	TGAGTACAAGTACTTGTGGGGGAGGGAATGGCACAGAGCAAAAGTTGAAGGGAAGGAAG	3420
Qy	3421	ATGAGAGGGCTCATGTTGTGGGGTGTGAAGGTCACTCCCTTTTCCATGTGATGGAGAT	3480
Dd	3421	ATGAGAGGGCTCATGTTGTGGGGTGTGAAGGTCACTCCCTTTTCCATGTGATGGAGAT	3480
Qy	3481	TAGAAAAACAGTGTGAGTTGTGATGCTTTCAGACACCCCCCACTATGAACATATCC	3540
Dd	3481	TAGAAAAACAGTGTGAGTTGTGATGCTTTCAGACACCCCCCACTATGAACATATCC	3540
Qy	3541	ACGAGGAGCGGACAGCTGTGGGAGACCTGGCATTTTAGGGAAGCGCGGCTTTTTCACAG	3600
Dd	3541	ACGAGGAGCGGACAGCTGTGGGAGACCTGGCATTTTAGGGAAGCGCGGCTTTTTCACAG	3600
Qy	3601	AGAACTTTATGCTCATCTCTTGTGCTACACTCCCACTTTGATGAGTTTCAGTCAAGT	3660
Dd	3601	AGAACTTTATGCTCATCTCTTGTGCTACACTCCCACTTTGATGAGTTTCAGTCAAGT	3660
Qy	3661	TTGCTTTCTACCGTTCTTGTGCTACACTGAGTGAAGTTCAGTGAAGTTCCTCCCAAGACGAGGA	3720
Dd	3661	TTGCTTTCTACCGTTCTTGTGCTACACTGAGTGAAGTTCAGTGAAGTTCCTCCCAAGACGAGGA	3720
Qy	3721	CAGCTCTTCTGTAAGGAGGACCTGATTTCAAGTGAAGTTCCTCCCAAGACGAGGA	3780
Dd	3721	CAGCTCTTCTGTAAGGAGGACCTGATTTCAAGTGAAGTTCCTCCCAAGACGAGGA	3780
Qy	3781	GAATCTAGGTCACAGTGAAGTTCAGTGAAGTTCCTCCCAAGACGAGGA	3840
Dd	3781	GAATCTAGGTCACAGTGAAGTTCAGTGAAGTTCCTCCCAAGACGAGGA	3840
Qy	3841	CCAGGTGAACGGTCACTGCTCAGATATCTGAGGTATTTGGGCTCCCAAGACGAGGA	3900
Dd	3841	CCAGGTGAACGGTCACTGCTCAGATATCTGAGGTATTTGGGCTCCCAAGACGAGGA	3900
Qy	3901	TCTGTTAGTGAAGTCACTGCTCAGATATCTGAGGTATTTGGGCTCCCAAGACGAGGA	3960
Dd	3901	TCTGTTAGTGAAGTCACTGCTCAGATATCTGAGGTATTTGGGCTCCCAAGACGAGGA	3960
Qy	3961	AAGAACTGTAAGGCTGAAGGAGACAGTGAAGGAGTATTTGGCAAGCCCAATACT	4020
Dd	3961	AAGAACTGTAAGGCTGAAGGAGACAGTGAAGGAGTATTTGGCAAGCCCAATACT	4020
Qy	4021	AAGCAATTCAGTGAAGGAGTGAAGGAGTATTTGGCAAGCCCAATACT	4080
Dd	4021	AAGCAATTCAGTGAAGGAGTGAAGGAGTATTTGGCAAGCCCAATACT	4080
Qy	4081	TACATTTTATTTGACTTGTCTACTATCTGGTCAATTTAGTGAAGTTCCTGAGCT	4140
Dd	4081	TACATTTTATTTGACTTGTCTACTATCTGGTCAATTTAGTGAAGTTCCTGAGCT	4140
Qy	4141	TAGTGGGCTCTATAGATCTTTCAATCTGCTCTAAATTTGTAAGTCAATTTCTGGAGCT	4200
Dd	4141	TAGTGGGCTCTATAGATCTTTCAATCTGCTCTAAATTTGTAAGTCAATTTCTGGAGCT	4200
Qy	4201	AGCAAAAGCTTACTGAGGAGTGAAGGAGTATTTGGCAAGCCCAATACT	4260
Dd	4201	AGCAAAAGCTTACTGAGGAGTGAAGGAGTATTTGGCAAGCCCAATACT	4260
Qy	4261	AGTCAATGCTTGAAGACAGATCCCTGATTTCCAGCTCTGCACTTGGCTAGTGGCCATGT	4320
Dd	4261	AGTCAATGCTTGAAGACAGATCCCTGATTTCCAGCTCTGCACTTGGCTAGTGGCCATGT	4320

Qy	4321	GTAATTAATTTGGCTTGAATTAAGTATTTGGGAAAGCCAGTTTCCACGACCTACATATC	4380
Dd	4321	GTAATTAATTTGGCTTGAATTAAGTATTTGGGAAAGCCAGTTTCCACGACCTACATATC	4380
Qy	4381	TGAAGAACCATTGATTTGAAAACCTAGAAAAGCTTGGGCAAAAATTTACTTAGAGATGTTTTTG	4440
Dd	4381	TGAAGAACCATTGATTTGAAAACCTAGAAAAGCTTGGGCAAAAATTTACTTAGAGATGTTTTTG	4440
Qy	4441	AGCTCATTTAAACCGGATGCTCTGAAAATGTTGGCAAAAATCAACCCAGAAATTAACAAAAGAG	4500
Dd	4441	AGCTCATTTAAACCGGATGCTCTGAAAATGTTGGCAAAAATCAACCCAGAAATTAACAAAAGAG	4500
Qy	4501	CTGGATTTGCAATATAGGACCAAGTATTTAGAAATCACTGGTATTAATAGCTATCATCTTAAT	4560
Dd	4501	CTGGATTTGCAATATAGGACCAAGTATTTAGAAATCACTGGTATTAATAGCTATCATCTTAAT	4560
Qy	4561	TAAAATATAGGGCTTATATATATTTAAGATTTAAACACAAAGAGTGGATAGCCCTCCCAAT	4620
Dd	4561	TAAAATATAGGGCTTATATATATTTAAGATTTAAACACAAAGAGTGGATAGCCCTCCCAAT	4620
Qy	4621	TTACTTTGGCTGCTTTTCAAAAGAGTAAAATATATCAGTCAATGGATTAATATAGTGTCTATG	4680
Dd	4621	TTACTTTGGCTGCTTTTCAAAAGAGTAAAATATATCAGTCAATGGATTAATATAGTGTCTATG	4680
Qy	4681	AAAGTATAGATCGAAACCCCTTTCTTACTTTTAACTTCACTTTCTTAACTTTTCTTCTT	4740
Dd	4681	AAAGTATAGATCGAAACCCCTTTCTTACTTTTAACTTCACTTTCTTAACTTTTCTTCTT	4740
Qy	4741	TTCAACCCCTGATCAAGCCACTAGTAAAGCACTATCTGCTGTGAGCTATTTATATGACTTT	4800
Dd	4741	TTCAACCCCTGATCAAGCCACTAGTAAAGCACTATCTGCTGTGAGCTATTTATATGACTTT	4800
Qy	4801	ACAGCAAAACAACTGCTGTGCTGCTTTTGGGAAAGGCAACAGGATAGCAGGAGCTC	4860
Dd	4801	ACAGCAAAACAACTGCTGTGCTGCTTTTGGGAAAGGCAACAGGATAGCAGGAGCTC	4860
Qy	4861	AGGCTAGCAAGTCTGACTTTGCCCTTAAAGCCAGAGGCAATGTTGATAGCAGAGAAAGTGA	4920
Dd	4861	AGGCTAGCAAGTCTGACTTTGCCCTTAAAGCCAGAGGCAATGTTGATAGCAGAGAAAGTGA	4920
Qy	4921	GCTCTTGGCAAGTGGGTGCTTAAAGTATCAGAAAACAGGAAAGGCTCCGGTGTGATGAAT	4980
Dd	4921	GCTCTTGGCAAGTGGGTGCTTAAAGTATCAGAAAACAGGAAAGGCTCCGGTGTGATGAAT	4980
Qy	4981	TATCAGTGAAGATATCTACCTCTTCTTCTATCGAACTTAAATCTCTCTTTCTTTCTTG	5040
Dd	4981	TATCAGTGAAGATATCTACCTCTTCTTCTATCGAACTTAAATCTCTCTTTCTTTCTTG	5040
Qy	5041	TGTGTAGGCTGATPAAACACACTTTGTTTCTTTTGGAGTGTTCATGGCTTTGTAGATTTTA	5100
Dd	5041	TGTGTAGGCTGATPAAACACACTTTGTTTCTTTTGGAGTGTTCATGGCTTTGTAGATTTTA	5100
Qy	5101	GTGCTCTGCCAGTCTTTGTAGAGGTTTGTACTTGTACCTTGACACCTGGCTGGATGTTAGC	5160
Dd	5101	GTGCTCTGCCAGTCTTTGTAGAGGTTTGTACTTGTACCTTGACACCTGGCTGGATGTTAGC	5160
Qy	5161	ATCCAAAAGGCAACACTTCTGAAATGCTGTGTAAAAGGTTATTTATTTACTTTTGTGTC	5220
Dd	5161	ATCCAAAAGGCAACACTTCTGAAATGCTGTGTAAAAGGTTATTTATTTACTTTTGTGTC	5220
Qy	5221	TTTGGAAAGGTGAAGGCTGTGTGAGAAAAGAACTCACAGGAGATGTGTTCTCTGTAGAAA	5280
Dd	5221	TTTGGAAAGGTGAAGGCTGTGTGAGAAAAGAACTCACAGGAGATGTGTTCTCTGTAGAAA	5280
Qy	5281	ACTTTTCTTCCCTTAAATGCTTATATTCCTTCTGTAAGGTTATTTATTTACTTTTGTGTC	5340
Dd	5281	ACTTTTCTTCCCTTAAATGCTTATATTCCTTCTGTAAGGTTATTTATTTACTTTTGTGTC	5340
Qy	5341	ATGCTGTCAATGAAGAGTGTGTTAGGCGGCTCTCATGCTCTGTGGAAAAGCACCATA	5400
Dd	5341	ATGCTGTCAATGAAGAGTGTGTTAGGCGGCTCTCATGCTCTGTGGAAAAGCACCATA	5400
Qy	5401	GGGAAAGGAATGTTATGCTGAGAAATCTGACCGGCAAGGAAATCTGGTTCAGAGCTCCCGG	5460

Db	5401	GGGAGGAATGTTATCTCGAATACTGACCGCAGGAAACTGGTCAGAGCTCCCCCG	5460
Qy	5461	AAGACCAACACAGGTGTTAAGTAGGAAACAGTCCAGGGTGGGCTCATGTAATAGAAATGGAA	5520
Db	5461	AAGACCAACACAGGTGTTAAGTAGGAAACAGTCCAGGGTGGGCTCATGTAATAGAAATGGAA	5520
Qy	5521	CAGAGCAGGGAAGATAGCTACAAAGTTCATAGGGTCCGGAGTCTTAAAGATACAAA	5580
Db	5521	CAGAGCAGGGAAGATAGCTACAAAGTTCATAGGGTCCGGAGTCTTAAAGATACAAA	5580
Qy	5581	TAGCTGCTTGGGCTTCATAACAAAGGAAGTCTGGAAGGAGGAGGAGGAGGAAATGG	5640
Db	5581	TAGCTGCTTGGGCTTCATAACAAAGGAAGTCTGGAAGGAGGAGGAGGAGGAAATGG	5640
Qy	5641	AAAGGGAAGAAACAGAAATAGTAGGAACTTGAAACAGCTACAAATCCTACACAGCAATTT	5700
Db	5641	AAAGGGAAGAAACAGAAATAGTAGGAACTTGAAACAGCTACAAATCCTACACAGCAATTT	5700
Qy	5701	TTCTTGGAAACAACTAGAGGTAGTAGGATAGGTGATGAGGGGAGCTTGCCTTGGCAT	5760
Db	5701	TTCTTGGAAACAACTAGAGGTAGTAGGATAGGTGATGAGGGGAGCTTGCCTTGGCAT	5760
Qy	5761	TTGAATCTGGGTTTTGTCTCTCCATTTGAGGTTGAAAGCGTCACCTTTTTACCCCTCGAA	5820
Db	5761	TTGAATCTGGGTTTTGTCTCTCCATTTGAGGTTGAAAGCGTCACCTTTTTACCCCTCGAA	5820
Qy	5821	TGAGAGGAAAGAGGGGTGTTATGACTCTCCTAGGAGTCTTACGTAGTTTACGCAATG	5880
Db	5821	TGAGAGGAAAGAGGGGTGTTATGACTCTCCTAGGAGTCTTACGTAGTTTACGCAATG	5880
Qy	5881	GAACAGACACTCGGGACCTCTCTTGACAAAAAATGGAACCTGTGTGTTCTTGTT	5940
Db	5881	GAACAGACACTCGGGACCTCTCTTGACAAAAAATGGAACCTGTGTGTTCTTGTT	5940
Qy	5941	TGTTCTTTTGTAAAGAACACAGGCAAGCCGACCAACATGGTTCGAATGCGGCTCTTT	6000
Db	5941	TGTTCTTTTGTAAAGAACACAGGCAAGCCGACCAACATGGTTCGAATGCGGCTCTTT	6000
Qy	6001	GAGTCAAGGGCTTTGAGTTAGGACTCATCAATAGTTGATCATGGTCAGGTGAGGGCTA	6060
Db	6001	GAGTCAAGGGCTTTGAGTTAGGACTCATCAATAGTTGATCATGGTCAGGTGAGGGCTA	6060
Qy	6061	CCTGTGAGGCGGACCTGCTGGCTTCGCACTTAAACATCTCCAGGTCTCAGTATCACTTC	6120
Db	6061	CCTGTGAGGCGGACCTGCTGGCTTCGCACTTAAACATCTCCAGGTCTCAGTATCACTTC	6120
Qy	6121	CTGCTACTTAGCACAGTTAGGAGTTGAGCAACCTTTTTTCCAAACCCCACTAAAAATTT	6180
Db	6121	CTGCTACTTAGCACAGTTAGGAGTTGAGCAACCTTTTTTCCAAACCCCACTAAAAATTT	6180
Qy	6181	AATTGACAAAAGACTGTGTAATTTGTGGGATACAGTGTGATAATGATCTATGTGTGCAT	6240
Db	6181	AATTGACAAAAGACTGTGTAATTTGTGGGATACAGTGTGATAATGATCTATGTGTGCAT	6240
Qy	6241	TGTCAGAGTTCAATAGATAGATTAATAGCCCATCAACAGCTTTATGGGTGGAATG	6300
Db	6241	TGTCAGAGTTCAATAGATAGATTAATAGCCCATCAACAGCTTTATGGGTGGAATG	6300
Qy	6301	CAAGTAATATAGGTAGATGCTGTGGTCTTCTTAGGTGAGAAAGGCAATGATTTTAAAGTTC	6360
Db	6301	CAAGTAATATAGGTAGATGCTGTGGTCTTCTTAGGTGAGAAAGGCAATGATTTTAAAGTTC	6360
Qy	6361	TTGGGCAAAATCATATATACCTATGCTATGCTATGCTATGCTATGCTATGCTATGCTATG	6420
Db	6361	TTGGGCAAAATCATATATACCTATGCTATGCTATGCTATGCTATGCTATGCTATGCTATG	6420
Qy	6421	AGAGGCTGATCTGTTGTTGCTCAGCAAGCAATGTCACAGCTCTTCTCACTAGT	6480
Db	6421	AGAGGCTGATCTGTTGTTGCTCAGCAAGCAATGTCACAGCTCTTCTCACTAGT	6480
Qy	6481	GTACCACTTTAGAAAAATGCTACCTGCTCAAAATGTTGTTGCTATTTCTATTTTTCATAGCT	6540

Db	6481	GTACCACTTTAGAAAAATGCTACCTGTGCTCAAAATGTTGTTGTTATCTTATTTTCATAGCT	6540
Qy	6541	TGAGAGAGTGGAGAGATCAAGCGGATTTGGGAACTGGACCTGCTGTTTATGCTCTGAG	6600
Db	6541	TGAGAGAGTGGAGAGATCAAGCGGATTTGGGAACTGGACCTGCTGTTTATGCTCTGAG	6600
Qy	6601	AAATGCTTGGCTCTGAGCGGAGAAAGCTAGAAAAAGAAAGAACTGCTCTTCTGCTTC	6660
Db	6601	AAATGCTTGGCTCTGAGCGGAGAAAGCTAGAAAAAGAAAGAACTGCTCTTCTGCTTC	6660
Qy	6661	TAAAAAGAAACAATTAAGATCCTCAATGGAATTTTAAAGAAAGTGAAGCTTAAC	6720
Db	6661	TAAAAAGAAACAATTAAGATCCTCAATGGAATTTTAAAGAAAGTGAAGCTTAAC	6720
Qy	6721	GTCCATCATCATTTAGAGATTTTCCACATGAAACCTGGCTCAGTTGAAAAAGAAATAGTGT	6780
Db	6721	GTCCATCATCATTTAGAGATTTTCCACATGAAACCTGGCTCAGTTGAAAAAGAAATAGTGT	6780
Qy	6781	CAAGTTGTCCATGAGACGAGGTAGACTTTGATAACCAAAAGATTCATTTGACAATATTT	6840
Db	6781	CAAGTTGTCCATGAGACGAGGTAGACTTTGATAACCAAAAGATTCATTTGACAATATTT	6840
Qy	6841	TATTGTCACTGATGATAACAGAAATAATGTACTTTTAAAAAATGTTTGAAGAGG	6900
Db	6841	TATTGTCACTGATGATAACAGAAATAATGTACTTTTAAAAAATGTTTGAAGAGG	6900
Qy	6901	TTACCTCTCATCTTCTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCATATTTT	6960
Db	6901	TTACCTCTCATCTTCTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCATATTTT	6960
Qy	6961	ATATATGTAAGTTTATTTTATTAAGTATACATTTTATTTATGTCAGTTTATTAATATGG	7020
Db	6961	ATATATGTAAGTTTATTTTATTAAGTATACATTTTATTTATGTCAGTTTATTAATATGG	7020
Qy	7021	ATTTATTTATAGAAAAATTTCTGCTATTTGATATTTAGTATAAGGCAAAATTAATTTATG	7080
Db	7021	ATTTATTTATAGAAAAATTTCTGCTATTTGATATTTAGTATAAGGCAAAATTAATTTATG	7080
Qy	7081	ACAATACTAGGAAACAGATATCTTAGGCTTTTAAACACATGATATCATATAATCT	7140
Db	7081	ACAATACTAGGAAACAGATATCTTAGGCTTTTAAACACATGATATCATATAATCT	7140
Qy	7141	TCGTCTTGTAAATTTTCTCCCTTTTAAATCAACATPACCATCATCATCATCATCATCAT	7200
Db	7141	TCGTCTTGTAAATTTTCTCCCTTTTAAATCAACATPACCATCATCATCATCATCATCAT	7200
Qy	7201	ATCATCTCATGATTTGATGCTTGAACCATATATATCTGTTAAAGTTGGTTCTGAGGCT	7260
Db	7201	ATCATCTCATGATTTGATGCTTGAACCATATATATCTGTTAAAGTTGGTTCTGAGGCT	7260
Qy	7261	CTGTGGTTTTGT	7320
Db	7261	CTGTGGTTTTGT	7320
Qy	7321	ATATTAGGTGCTTCTCTCATCATGCTTTGCTCTTATTTATTTAGACAGAGGCTCTGCTCTG	7380
Db	7321	ATATTAGGTGCTTCTCTCATCATGCTTTGCTCTTATTTATTTAGACAGAGGCTCTGCTCTG	7380
Qy	7381	AACTGTAGCTAGGCTGGCCAAAGCTCTATTAATTTTTTTTTTAAAGATTAATTAATTTATG	7440
Db	7381	AACTGTAGCTAGGCTGGCCAAAGCTCTATTAATTTTTTTTTTAAAGATTAATTAATTTATG	7440
Qy	7441	TGTAT 7445	
Db	7441	TGTAT 7445	

RESULT 3  
 US-09-751-797-29  
 ; Sequence 29, Application US/09751797  
 ; Patent NO. US20010024652A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Dumoutier, Laure



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; APPLICANT: Louhed, Jamila
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors
; TITLE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof
; FILE REFERENCES: LUD 5543.2
; CURRENT APPLICATION NUMBER: US/09/751.797
; CURRENT FILING DATE: 2000-12-29
; PRIOR APPLICATION NUMBER: 09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 29
; LENGTH: 5935
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-09-751-797-29

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Query Match		57.0%;	Score 4245.2;	DB 3;	Length 5935;
Best Local Similarity		88.3%;	Pred. No. 0;		
Matches 5039;		Conservative	0;	Mismatches 178;	Indels 487;
Gaps		20;			
QY	1971	TACATGCTACCGACGAA	CATGCTCCCTGATGTTTTTGGCTTTTGCTCTCTCACTAAC	2030	
DB	293	TACATGCTATCCGACGAGATG	TCCCTGATGTTTTTGGCTTTTGCTCTCTCGCTAAC	352	
QY	2031	AGGCTCTCCTCTCACTTATCAACTTGTTGACACTTGTGCGATCTCTGATGCTGTCCTGCA	2090		
DB	353	AGGCTCTCCTCTCAGTTATCACTTTTGTACACTTGTGCGATCGGTATGCTGTCCTGCA	412		
QY	2091	GAATCTATGAGTTTTTCCCTTATGGGAC	CTTTGGCCGACGTGCTTCTCAATGC	2150	
DB	413	GAATCTATGAGTTTTTCCCTTATGGGAC	CTTTGGCCGACGTGCTTCTCAATGC	472	
QY	2151	CTGTGGGCCCGAGGACAAATGGCTGCGCGTCAACACCGGTGCAAGCTTGAGGTGC	2210		
DB	473	CTGTGGGCCCGAGGACAAATGGCTGCGCCATCAACACCGGTGCAAGCTTGAGGTGC	532		
QY	2211	CAACTTCCAGCAGCCGTAATCGTCAACCGCACCTTTATGCTGCGCAAGGAGGTACAGCT	2270		
DB	533	CAACTTCCAGCAGCCGTAATCGTCAACCGCACCTTTATGCTGCGCAAGGAGGTACAGCT	592		
QY	2271	GCATCTCTTTCTCTCCATACGCGCTTGCCCATTTTTCTGGAAGCACTTGCAAACTCTTTAG	2330		
DB	593	GCATCTCTTTCTCTCCATACGCGCTTGCCA - TTTCTCTGAAGCACCTTGCAACTCTTTAG	651		
QY	2331	GGCGCTTTATCTCGCAGGTCTCACTACCTATGTTTTTCTGTTTATGAGACTCTTTTA	2390		
DB	652	GGCGCTTTATCTCGCAGGTCTCACTACCTATGTTTTTCTGTTTATGAGACTCTTTTA	711		
QY	2391	AGGACTGGGTCTTTTCTATTTCTATTTCAAGGTCTCAGGACCATTTTCCCTCTTGGCCT	2450		
DB	712	AGGACTGGATCTTTTCTATTTCTATTTCAAGGTCTCAGGACCATTTTCCCTCTTGGCCT	771		
QY	2451	TCAGGACACATATACTGAA'TTTTTATCTACAGAGCGCATTTAGAAAGCCACCCAGCACTG	2510		
DB	772	TCAGGACACATATACTGAA'TTTTTATCTACAGAGCGGTTTAGAAAGCCACCCAGCACTG	831		
QY	2511	CAATACTTTTCCATTTCTCTGTCCTCTTCTGAACTCATCTCTTTGGCTACTCTCTGAG	2570		
DB	832	CAATACTTTTCCATCTCTGTCCTCTTCTGAACTCATCTCTTTGGCTACTCTCTGAG	891		
QY	2571	ACCCACTGCGGACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTTGTCAACCCAGG	2630		
DB	892	ACCCACTGCGGACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTTGTCAACCCAGG	951		
QY	2631	CACTTAGGTTTTTCTCTCTTTTCAGGCCAGCCTTGCAATAAACAACAGACGTCCGGCTC	2690		
DB	952	CACTTAGGTTTTTCTCTCTTTTCAGGCCAGCCTTGCAATAAACAACAGACGTCCGGCTC	1011		
QY	2691	ATCGGGGAGAACTGTTTCCGAGGAGTCAGTGTGAAGTCCTCACTGTGATGAGCGGCTAG	2750		

1012	ATCGGGAGAACTGTTCCGAGAGTCAGTGTAAAGTCCTCACTGTGATGACGAGGCGTAG	107
2751	CTCGGGAGCTGGTGGACCCCTCTGGGATAGTCTGACGTATGACCCCTGCTGCTTCTTGTC	2810
1072	CTCGGGAGCTGGTGGACCCCTCTGGGATAGTCTGACGTATGACCCCTGCTGCTTCTTGTC	1131
2811	TACCTGCAGGCTAAAGATCAGTCTGCTACCTGATGAAGCAGGTGCTCAACTTCACCCCTGGAA	2870
1132	TACCTGCAGGCTAAAGATCAGTCTGCTACCTGATGAAGCAGGTGCTCAACTTCACCCCTGGAA	1191
2871	GAGTTCCTGCTCCCCAGTCAGACAGCTTCCAGCCCTACATGACGAGGTGGTACCTTTTC	2930
1192	GAAITCTGCTCCCCAGTCAGACAGGTTCCGGCCCTCATGACGAGGTGGTGGCCTTTC	1251
2931	CTGACCAAACTCAGCAAACTCAGCTCAGCTCCTGTGTAAAGTCTGACTCTGGCTACCTATGCT	2990
1252	CTGACCAAACTCAGCAAACTCAGCTCAGCTCCTGTGTAAAGTCTGCTCTGGCTACCTATGCT	1311
2991	CCTCTCTCTTCTCTTCTATTCAGTAAGAAACCGGAGGTCTCGGCTCTCTCTCTTCACA	3050
1312	CCTCTCTCTTCTCTTCTATTCAGTAAGAAACCGGAGGTCTCGGCTCTCTCTCTTCACA	1371
3051	AGAGTGAGGAGGCGCTCAGCACCAACCAACCATATAGCCCACTTGAATAGTTCACAAAGG	3110
1372	AGAGTGAGGAGGCGCTCAGCACCAACCAACCATATAGCCCACTTGAATAGTTCACAAAGG	1431
3111	CTTTGGCTTCAAATTGAGTAATACTTTTGAGTTTCTGTATGAGTGAAGCTTTATTTGTTTTATC	3170
1432	CTTTGGCTTCAAATTGAGTAATACTTTTGAGTTTCTGTATGAGTGAAGCTTTATTTGTTTTATC	1491
3171	CATGGAAAGAAATCAACTCAAATCTGTAGGTGAGAAAGATGTTGGGAACGAAAAAAGG	3230
1492	CATGGAAAGAAATCAACTCAAATCTGTAGGTGAGAAAGATGTTGGGAACGAAAAAAGG	1551
3231	CCTAGATAGAGAAACAGACTCTGCTGAGTATAGTACTTTAT ---GGGGGAGCAGGGGGCG	3286
1552	CCTAGATAGAGAAACAGACTCTGCTGAGTACAGTACTTTATGGGGGGGGGGGAGGGGGCG	1611
3287	ATATCCACTGAGTACAAGTACTTTGTGGGAGAGAAATCCACTGAGTACAAAGTACTTGTG	3346
1612	ATATCCACTGAGTCCAAGTACTTTGTGGGAGAGAAATCCACTGAGTACAAAGTACTTGT--	1669
3347	GCATGGAGATCCACTGAGTACAAGTACTTTGTGGGGGAGGGAATGGCACAGACAAAAGT	3406
1670	-----GGGGGAGGAATGGCACAGACAAAAGT	1697
3407	TGAAGGGA---ACGAAATGGAGAGCGCTCATGGTTGGGGGTGTGAAGTCACTCC--TT	3462
1698	TGAAGGAAAGAGAGAGATGGAGAGCGCTCAATGTGTTGGGGGTGTGAAGTCACTCCCTTT	1757
3463	TTCCATGTGATGAGAGTTAAGAAAAACCAAGTGTGTGAGTTTCATGTCTTTTCAGACACCCC	3522
1758	TTCCATGTGATGAGAGTTAAGAAAAATCAGTGTGTGAGTTTGATGTCTTCAGACACCCC	1817
3523	CAACTATGMAACATATCCACGAGGAGCGGGCAGACTGTGGGAGACCTGGCACTTTAGGGAA	3582
1818	AA-----CTATGSCACACTGTGGGAGACCTGGCACTTTAGGGAA	1855
3583	GGCGCGGCTTTTACACGAGAAACTTTATGCTCATCTCTTTGTGCTACATCCCACTTTG	3642
1856	GGCGCGGCTTTTACACGAGAAACTTTATGCTCATCTCTTTGTGCTACATCCCACTTTG	1915
3643	ATGAGTTACGCTCAGGTTTTGTTTTCTACCGTTCTTGTCTACTCGTGAAACTTCAGTAGG	3702
1916	ATGAGTTAAGCTCAGGTTTCGTTTTCTACCGTTCTTGTCTACTCGTGAAACTTCAGTAGG	1975
3703	ATTCCCCAAAGACGAGACAGCTCTTCTGTGAAGGAGGGAACCTTGGATTTTCAGTGTCTCTAG	3762
1976	ATTCCCCAAAGACGAGACAGCTCTTCTGTGAAGGAGGGAACCTTGGATTTTCAGTGTCTCTAG	2035
3763	AGAACGAAATAGCTCAGAGAACTTAGGTCAAACGTGAAATCTAGGTCAACGCGGCAAAA	3822
2036	AGAACGAAATAGCTCAGAGAACTTAGGTCAAACGTGAAATCTAGGTCAACGCGGCAAAA	2095

Qy	3823	TGACTGAACGCCCTCTATTTCAGAGTGAAAGCGTCAAGTGCCTCAGATATACAGGTATTGG	3882
Db	2096	TGACTGAACGCCCTCTATTTCAGAGTGAAAGCGTCAAGTGCCTCAGATATACAGGTATTGG	2155
Qy	3883	GCTCCCAACCGGATAAGATTCTGTGTAGTGAAGTCTGCTTTTATTTTTCAGACATCAGCGGT	3942
Db	2156	GCTCCCAACCGGATAAGATTCTGTGTAGTGAAGTCTGCTTTTATTTTTCAGACATCAGGTGT	2215
Qy	3943	GACGACAGAACATCCAGAGAAATGTGAGAAAGGCTGAAGGAGACAGTGAAGAGTACTA	4002
Db	2216	GACGACAGAACATCCAGAGAAATGTGAGAAAGGCTGAAGGAGACAGTGAAGAGTACTA	2275
Qy	4003	TTGCAAGGCCACAATACTAAGGCCATTCAAGTAGGAGAGTGGGAGATTCTTTCTCTGCTTC	4062
Db	2276	TTGCAAGGCCACAATACTAAGGCCATTCAAGTAGGAGAGTGGGAGATTCTTTCTCTGCTTC	2335
Qy	4063	CCAGTCCCTTCTACTTTTGTGAACATTTTATTTGACTTGTCTACTATCTGGTCCATTAATCG	4122
Db	2336	CCAGTCTCTTCTACTTTTGTGAACATTTTCTTTGACTTGTCTACTTGTCTGGTCCATTAATCG	2395
Qy	4123	CTTAGCTGCACTGTATCTAGCTGGGTCTATAGATCTTTTCAATCTGTGTCTAAATTTGTA	4182
Db	2396	CTTAGCTGCACTGTATCTAGCTGGGTCTATAGATCTTTTCAATCTGTGTCTAAATTTGTA	2455
Qy	4183	AGTCACAAATCTGAGCTAGCAGAAAGCTTAGCTCAGCAGTCTCATGAGCACTTGCTCG	4242
Db	2456	AGTCACAAATCTGAGCTAGCAGAAAGCTTAGCTCAGCAGTCTCATGAGCACTTGCTCG	2515
Qy	4243	GAGGATGGCTTGTGACAGAGTCAATGTCTAGAAAGACAGCATCCCTGATTTCCAGCTTGCA	4302
Db	2516	GAGGATGGCTTGTGACAGAGTCAATGTCTAGAAAGACAGCATCCCTGATTTCCAGCTTGCA	2575
Qy	4303	CTTGCCCTAGTGCCCATGTGTAAATTTACTTTTGGCTTGAATTAAGTATTTTGGAAAGCCAGTTC	4362
Db	2576	CTTGCCCTAGTGCCCATGTGTAAATTTACTTTTGGCTTGAATTAAGTATTTTGGAAAGCCAAATTC	2635
Qy	4363	CCACGGACCTACATAATCTGAAGAACCAATGATTTGAAACCTTAGAAAGCTGGCACAAACT	4422
Db	2636	CCACGGACCTACATAATCCGAAAGAGCATGCAITTTGAAACCTTAGAAAGCTGGCACAAACT	2695
Qy	4423	TACTAGAGATGATTTTGGAGTCAATTAACCGGATGCTCTGAAATGTGGCAAAATCAACCC	4482
Db	2696	TACTAGAGATGATTTTGGAGTCAATTAACCGGATGCTCTGAAATGTGATCAAAATCAACCC	2755
Qy	4483	AGAAATACAAACAAAGAGCTGGATTGGAAATAGGACAAAGTATTTAGATCACTGGTATT	4542
Db	2756	AGAAATACAAACAAAGAGCTGGATTGGCAATAGGACAAAGTATTTAGAAATCACTGGTATT	2815
Qy	4543	AATAGCTATCATCTTAATTAATAATATAGGCTATATA - - - TATATTTAAGATTAAACA	4598
Db	2816	AACAGCTGTATCTTAATTAATAATATAGTGTATTTAGCTGGCTATTTAAGATTAAACA	2875
Qy	4599	CAAGAGTGGATAGCTCCCAATTTACTTTGGCTCGTGTTCCTGAAAGAGTAAATAATCAGTC	4658
Db	2876	CAAGAGTGGATAACTTCCCAATTTACTTTGGCTCGTGTTCCTGAAAGAGTAAATAATCAGTC	2935
Qy	4659	ATGGAATTAATATAGTGTCAAGAAAGTATAGATGGAAACCCCTTTCCTTACCTTTTACCT	4718
Db	2936	ATGGAATTAATATAGTGTCAAGAAAGTATAGATGGAAACCCCTTTCCTTACCTTTTACCT	2995
Qy	4719	TCATTTCTTACT - - - TTTTCTTCTTCTACCCCTGATCAAGCCACTAGTAAGCACT	4773
Db	2996	TCATTTCTTACTTATTTTCTTCTTCTACCCCTGATCAAGCCACTAGTAAGCACT	3055
Qy	4774	ATCTGCTGAGCTATTATATGACTTTACAGCAACAAACATTTGCTGTGGTCCCTTTTGG	4833
Db	3056	ATCTGCTGAGCTATTATATGACTTTACAGCAACAAACATTTGCTGTGGTCCCTTTTGG	3115
Qy	4834	GGAAAGGAAAGGATAGCAGGAGGCTCAGGCTAGCAAGTCT - GACTTGGCCCTTAAGCCAG	4892
Db	3116	GGAAAGGAAAGGATAGCAGGAGGCTCAGGCTAGCAAGTCTCAGGCTCAACCTCAAGCCAG	3175

Qy	4893	AGGCATGCTTTGATAGCAGAGAAAGTGAAGCTCTTCGCAAGTGGGTGCTTTAAGTAATCA	4952
Db	3176	AGGCATGCTTTGATAGCAGAGAAAGTGAAGCTCTTCACAAGTGGGTGCTTTAAGTAATCA	3235
Qy	4953	GAAACAGGAAGGCTCCGGTTGATGGAAATTAATCAAGTAAGATATCTACCCCTTATCTC - - - CT	5009
Db	3236	GAAACAGGAAGGCTCCGGTTGATGGAAATTAATCAAGTAAGATATCTACCCCTTATCTCCTCT	3295
Qy	5010	TCATATCGAACTAAATCGTCTCTTTTCTTGTGTGTAGGCTGATAAACAACACTTGTGTTTC	5069
Db	3296	TCATATAGAACTAAACCGTCTCTCTCTTGTGTGTAGGCTGATAAACAACACTTGTGTTTC	3355
Qy	5070	TTTTGAGTGTTCATGGCTTTTGTAGATTTTATGTGCTCTGCCAGTCTTGTGTAGAGGTTT	5129
Db	3356	TTTTGAGTGTTCATGGCTTTTGTAGATTTTATGTGCTCTGCCAGTCTTGTGTAGAGGTTT	3415
Qy	5130	GTTTACCTTGAACACTGGGCTTGGATGTAGCATGCCAAAGGCACACACTTCTGNAATGCT	5189
Db	3416	GTTTACCTTGAACACTGGGCTTGGATGTAGCATGCCAAAGGCACACACTTCTGNAATGCT	3475
Qy	5190	GTGTAAAAGGTATTATTCAATTTTACTTTTGTGTTGAAAGGTGAAGCTGTGTGAGAAAG	5249
Db	3476	GTGTAAAAGGTATTATTCAATTTTACTTTTGTGTTGAAAGGTGAAGTGTGTGAGAAAG	3535
Qy	5250	AACTCACAGGAGATGTCTCTGTAGGAAACCTTTTTTTTTCCCTTTAAATGCTATAA	5309
Db	3536	AACTCACAGGAGATGTATTCTGTAGGAAAC - TTTTTTTTTCCCTTTAAAGCTATAA	3594
Qy	5310	TCACATTTTCACTTGTGACTTTTATACATGCTGTACATGAAAGAGTGTGTAGGCC	5369
Db	3595	TCACATTTTCACTTGTGACTTTTATACATGCTGTACATGAAAGAGTGTGTAGGCC	3654
Qy	5370	CGCTCTCATCGCTCTGGGAAAGCACCACATAGGGGAAGGAATGTTATGCTGAGAAATCTG	5429
Db	3655	CGCTCTCGTGGCTCTGGGAAAGCACCATAGGGGAAGGAATGTTATGCCGAGAAATCTG	3714
Qy	5430	ACCGGACGGGAAACTGGTCTAGAGCTCCCCGGAAGACACACAGGTGTGTAAGTAGGAACA	5489
Db	3715	ACTGGCAGGGAACCTGGTCTAGAGCTCCCCAAGACCACTACAGGTGTGTAAGTAGGAACA	3774
Qy	5490	GTCAGAGGTGGGTCTATGTAATAGAAATGGAAACAGAGCGAGGGAAGATAGCTACAAAGTT	5549
Db	3775	GTCAGAGGTGGGTCTATGTAATAGAAATGGAAACAGAGCGAGGGAAGATAGCTACAAAGTT	3834
Qy	5550	TCATAGGCTCCGAGTCTTAAAGATACAAATAGCTCTTGGGCTTCATAACAAGGAAG	5609
Db	3835	TCATAGGCTCCTAAGTCTTTAAGATACAAATAGCTGTGGGCTTCATAACAAGGAAG	3894
Qy	5610	TCTGGGAAGGCAGCAAG - - - TGAGAGGGAATGGAAAGGGAAGGGAAGGAGTGTAGGGA	5666
Db	3895	TCTGGGAAGGCAGCAAGCAATTTGAGAGGAGATGGAAAGGGAAGGGAAGGAGTGTAGGGA	3952
Qy	5667	CTTGAACAGCTACAAATCTCTACACAGATTTTTCTTGGAAACAATCTAGAA - - - GGT	5722
Db	3953	TTTGAAGAGCTACAAATCTCTACACAGAGGATTTTTCTTGGAGGAATCTAGAAACAAGGT	4012
Qy	5723	AGTGGATTTAGGT - GATTTGACAGGGGACTTGTCTTGGCAATTTGAATCTGGGTTTTGTCTC	5781
Db	4013	GGTGGATTTAGGTGATCGCAGAGGACTTGTCTTGGCAATTTGAATCTGGGTTTTGTCTC	4072
Qy	5782	TCCATTGAGGTGGAAGCGTCACCC - TTTTTTACCTCGAAATGGAGGGAAGGAGGAGTGT	5840
Db	4073	TCCATTGAGGTGAGAGGTCACCCCTTTTTTACCTCGATAGGAGGGAAGGAGGAGTGT	4132
Qy	5841	GTTATGATCTCTACTGAGGTTTTTACTAGTTTACGCAATGGAAACAGACACTCGGGACCTC	5900
Db	4133	GTTTGTGATCTCTACTGAGGTTTTTACTAGTTTACGCAATGGAAACAGACACTCGGGACCTC	4192
Qy	5901	CTCTTTGAC - - - - - AAAAAAATGGAAACCTGTTGTTGTTGTTGTTGTTGTTGTTG	5950
Db	4193	CTCTTTGACAAAAAATAAAAAAAGAAACCTGTTGTTGTTGTTGTTGTTGTTGTTG	4252
Qy	5951	TTAAGAAAGCAC - - - - -	5963

Db	4253		TTAAGAAGCACAGCGAGCTGGGCATGGTGGCCCATGCCCTTAAATCCACGATTTGGGAG	4312
Qy	5964	-----	-----	5963
Db	4313	GCAGAGCGAGTGACTTCTTAATTTCAAGGCCAGCCTGGTCTACAAAGTGAGTTCAGGA	4372	
Qy	5964	-----	-----	5963
Db	4373	CAGCCAGGGCTATACAGAGAAACCTGTCTCGGAAAAAAGAAAGAAAGAAAG	4432	
Qy	5964	-----	-----	5963
Db	4433	AAAAGAGAGAGAGGAGAGAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG	4492	
Qy	5964	-----	-----	5963
Db	4493	GAGAGGAGAGGAA	4552	
Qy	5964	-----	-----	5963
Db	4553	GAA	4612	
Qy	5964	-----	GGCAAGCCGACCACTGGGT	5985
Db	4613	AAGAGAAAAAGAAAAAGCAAGCAAGCACTGGCAAGCATGCCACATCGGGA	4672	
Qy	5986	TGAATGTGGGTCTTTGAGTCAAGGCTTTTGAGTTGAGCACTCATCAATAGTTGATCATGG	6045	
Db	4673	CGTATGTGGGTCTTTGAGCAGAGGCTTTGAAATGAGCGCTCATCAATAGTTGATCATGG	4732	
Qy	6046	TCAGGTGGAGGGCTACCTGTGAGSCGAGCCCTGCTGGCTTCGCACCTTAAACATCTCCAGG	6105	
Db	4733	TCAGGTGGAGGGCTACCTGTGAGSCGAGCCCTGCTGGCTTCGCACCTTAAACATCTCCAGG	4792	
Qy	6106	TCTCAGTATCACTTCCTGCTACTTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTCCAA	6165	
Db	4793	TCTCAGTATCACTTCCTGCTGCTTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTCCAA	4852	
Qy	6166	CCCCACTAAAAATTAATGCAAAAAACACTGTGTAAATTCGTGGGATACAGTGTGATAAT	6225	
Db	4853	CCCCACTAAAAATTAATTAAGAGGCAAGTAAATTTGTTGGGATACAGTGTGATAAT	4912	
Qy	6226	GATCTATGTGTGCAATGTGCAAGGTTCAATAAGATAGATTAATAGGCCCATCAACAGCTT	6285	
Db	4913	GATCTATGTGTGCAATGTGCAAGGTTCAATAAGGATAGATCAATAGGCCCATCAACAGCTT	4972	
Qy	6286	TATGGGTGTGAATGCAAGTAAATATAGGTATAGTCCTGTGGTGTCTTATGGTCAGAAAGG	6345	
Db	4973	TATGGGTGTGAATGCAAGTAAATATAGGTATAGTCCTGTGTGTCTTATGGTCAGAAAGG	5031	
Qy	6346	CATGATTTTAAGGCTTTGGGCAATCATATTAATCACTCATGCTAAAAATACATTAATGTGA	6405	
Db	5032	CATGATTTTAAGGCTTTGGGCAATCATATTAATCACTCATGCTAAAAATGATTAATGTGA	5091	
Qy	6406	TTATTAATCTTTTAGAGAAGCGTGATCTTGGTTTTGGTGCTCAGCAAGCAAAATGTCAAC	6465	
Db	5092	TTATCAATCTTTTAGAGAAGCGTGATCTTGGTTTTGGTGCTCAGCAAGCAAAATGTCAAC	5151	
Qy	6466	AGCTCTTTCTAACTGGGTACCACTTTAGAAAAATGCTCTGTGCTCAAAATGGTTGTATT	6525	
Db	5152	AGCTCTTTCTAACTAGTACCACCTTTAGAAAAATGCTACCCGTGCTCAAAATGGTTGTATT	5211	
Qy	6526	CTTAATTTTCATAGCTCGGAGAGTGAGAGATCAAGCGATTTGGGAACTGACCTGCT	6585	
Db	5212	CTTAATTTTCATAGCTCGGAGAGCGGAGAGATCAAGCGATCGGGGAACCTGACCTGCT	5271	
Qy	6586	GTTTATGTCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAAACGAGAACTG	6645	
Db	5272	GTTTATGTCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAAACGAGAACTG	5331	
Qy	6646	CTCCTTCCTCCCTTCTAAAAAGAAACAATAAGATCCCTGAATGGACCTTTTATCTAAAGGA	6705	

PRECIT.T 4

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US-10-627-273-29
; Sequence 29, Application US/10627273
; Publication No. US20040110189A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid
; TITLE OF INVENTION: (Title) The Proteins
; FILE REFERENCE: LUD 5543.2
; CURRENT APPLICATION NUMBER: US/10/627,273
; CURRENT FILING DATE: 2003-07-25
; PRIOR APPLICATION NUMBER: US/09/751,797
; PRIOR FILING DATE: 2000-12-29
; PRIOR APPLICATION NUMBER: 09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US/09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 29
; LENGTH: 5935
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-10-627-273-29

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Query Match 57.0%; Score 4245.2; DB 7; Length 5935;  
Best Local Similarity 88.3%; Pred. No. 0;  
Matches 5039; Conservative 0; Mismatches 178; Indels 487; Gaps 20;

Db	5332	CTCCTTCCTCGCCTTCTTAAAAAGAACAAATAAGATCCCTCGAATGGACCTTTTTTACTATAAGGA	5391
Qy	6706	AAGTGAAGAGCTAACGCTCCATCATCATTAGAAGATTTACATGAAACCTGGCTCAGTTGA	6765
Db	5392	AAGTGAAGAGCTAACGCTCCACCATCATTAGAAGATTTACATGAAACCTGGCTCAGTTGA	5451
Qy	6766	AAAAGAAAAATAGTGTCAAGTTGTCCATGAGACACGAGGTTAGACTTGTGATAACCAAAAGAT	6825
Db	5452	AAGAGAAAAATAGTGTCAAGTTGTCCATGAGACACGAGGTTAGACTTGTGATAACCAAAAGAT	5511
Qy	6826	TCATTGACAATATTTTTATCTGCTACTGATGATACAAAGAAAAATAATGTACTTTTAAAAAAA	6885
Db	5512	TCATTGACAATATTTTTATCTGCTACTGATGATGACAAAGAAAAAGATGTACTTTTAAAAAAA	5571
Qy	6886	TTGTTTGAAGAGGTTACCTCTCATTCCTTTTAGAAAAAAGCTTATGTAACTTCATTTC	6945
Db	5572	TTGTTTGAAGAGGTTACCTCTCATTCCTCTAGAGAAAAAGCCTATGTAACTTCATTTC	5631
Qy	6946	CATATCCAATATTTTATATATGTAAGTTTATTTATTAAGTATACACATTTTATTTATGTGC	7005
Db	5632	CATAACCAATACCTTTATATATGTAAGTTTATTTATTAAGTATACACATTTTATTTATGTGC	5691
Qy	7006	AGTTTATTAATATGGATTTATTTATAGAAACATTCCTGCTATTGATATTT - AGTATAAG	7064
Db	5692	AGTTTATTAATATGGATTTATTTATAGAAAAATTAUCTGATGTTGATATTGAGTATAAA	5751
Qy	7065	GCAAAATAATATTATGACAAATACTATGGAAAAAAGATATCTTAGGCTTTTAATAAACACA	7124
Db	5752	GCAAAATAATATTATGATAATAACTATAGAAAAAAGATATCTTAGGCTTTTAATAAACACA	5811
Qy	7125	TGGATATCAAAATCTTCGCTGTGTAATTTTTTCTCCCTTTTAATATACAACAATACCATCA	7184
Db	5812	TGAATATCAAAATCTTCGCTGTGTAATTTTTTCTCCCTTTTAATATACAACAATACCATCA	5871
Qy	7185	TCATCATCATTAACCAATCATTTCTCATGATTTTCATGCTTGACCCCATATTATACGTGTAAA	7244
Db	5872	TCGTCATCATTAACCAATCATTTCTCATGACTTCATGCTTGACTCATATTATCTGTTAAG	5931
Qy	7245	GTTCG 7248	
Db	5932	TTTCG 5935	

Query Match 57.0%: Score 4245.2: DB 7: Length 5935:

Query Match      37.0%; Score 4243.2  
Best Local Similarity      88.3%; Pred. No. 0;

**Matches 5039; Conservative 0; Mismatches 178; Indels 487; Gaps 20;**

Qy	1971	TACCATGCTACCGACGAA	CATGCTCCCTGTAGTGT	TTTTCGCTTTTTCGCTCTCTCACTAAC	2030
Dd	293	TACCATGCTATCCGACGAG	ATGTTCCCTGTAGTGT	TTTTCGCTTTTTCGCTCTCTCGCTAAC	352
Qy	2031	AGGCTCTCCCTCTCACTT	ATCAACTGTGTGACACT	TTGTGCGCATCTCTGTAGTGGCTGTCTCTGCA	2090
Dd	353	AGGCTCTCCCTCTCACTT	ATCAACTTTTGTACACT	TTGTGCGCATCTCTGTAGTGGCTGTCTCTGCA	412
Qy	2091	GAATCTATGAGTTTTTCC	CTTATGCGGACCTTTGG	CGCGCAGAGTCCCTCTCTCATATGC	2150
Dd	413	GAATCTATGAGTTTTTCC	CTTATGCGGACCTTTGG	CGCGCAGAGTCCCTCTCTCATATGC	472
Qy	2151	CCTGTGGCGCCAGAGG	CAAAATGCGTCCCGTCA	ACACCGGTGCAAGCTTGAAGTGC	2210
Dd	473	CCTGTGGCGCCAGAGG	CAAAATGCGTCCCGTCA	ACACCGGTGCAAGCTTGAAGTGC	532
Qy	2211	CAACTTCAGCAGCGGT	ACATCGTCAACCGCAC	CTTTATGCTGGCCAAAGGAGGTACAGCT	2270
Dd	533	CAACTTCAGCAGCGGT	ACATCGTCAACCGCAC	CTTTATGCTGGCCAAAGGAGGTACAGCT	592
Qy	2271	GCATCTCTTCTCCAN	ACGCGCTTGCCATTTT	CTGGAAGCACTTGCAAACTCTTTTAG	2330
Dd	593	GCATCTCTTCTCCAN	ACGCGCTTGCCCA--	TTTCTCTGAAGCACTTGCAAACTCTTTTAG	651
Qy	2331	GGGCGCTTATCTCCG	CAGGTCCTCACTATG	TTTCTCTCTTTAGAGACTCTTTA	2390
Dd	652	GGGCGCTTATCTCCG	CAGGTCCTCACTATG	TTTCTCTCTTTAGAGACTCTTTA	711
Qy	2391	AGGACTGGGCTTTTCT	ATTTCTATTTCAAGG	TCCTCAGGACCACTTTCTATCTTGGCCT	2450
Dd	712	AGGACTGGGCTTTTCT	ATTTCTATTTCAAGG	TCCTCAGGACCACTTTCTATCTTGGCCT	771
Qy	2451	TCAGGACACATATCT	GAATTTTATCTACG	AGGCGCATTTTGAAGCCACCCACGACTG	2510
Dd	772	TCAGGACACATATCT	GAATTTTATCTACG	AGGCGCATTTTGAAGCCACCCACGACTG	831
Qy	2511	CAATACCTTCCATTT	CTCTGTGCTCTCTCT	GTGAATCATCTCTCTTGGCTACTCTCTGAG	2570
Dd	832	CAATACCTTCCATTT	CTCTGTGCTCTCTCT	GTGAATCATCTCTCTTGGCTACTCTCTGAG	891
Qy	2571	ACCCACTCGCGGACA	TACATCTCTACTTAC	AGGCTTTTCTTCCATCTCTCTGTCAACCCAGG	2630
Dd	892	ACCCACTCGCGGACA	TACATCTCTACTTAC	AGGCTTTTCTTCCATCTCTCTGTCAACCCAGG	951
Qy	2631	CACCTTAGGGTTTTCT	CTTTTCAGGCGAGCT	TCGAGATTAACAACAACAGAGTCCGGCTC	2690
Dd	952	CACCTTAGGGTTTTCT	CTTTTCAGGCGAGCT	TCGAGATTAACAACAACAACAGAGTCCGGCTC	1011
Qy	2691	ATCGGGGAGAACTGT	TCCGAGGAGTCAGT	GAAGTCTCTCACTGTGATGAGCGGGCTAG	2750
Dd	1012	ATCGGGGAGAACTGT	TCCGAGGAGTCAGT	GAAGTCTCTCACTGTGATGAGCGGGCTAG	1071
Qy	2751	CTCGGGGAGCTGTGG	ACCTCTGGGATAGT	CTGACAGTATGACCCCTGCTCTTCTGTC	2810
Dd	1072	CTCGGGGAGCTGTGG	ACCTCTGGGATAGT	CTGACAGTATGACCCCTGCTCTTCTGTC	1131
Qy	2811	TACCTGACGGTAAAG	ATCAGTGTACCTGT	ATGAAGAGAGTGTCTCAACTCACCCCTGGAA	2870
Dd	1132	TACCTGACGGTAAAG	ATCAGTGTACCTGT	ATGAAGAGAGTGTCTCAACTCACCCCTGGAA	1191
Qy	2871	GACGTTCTGCTCC	CCAGTCAGAGGTTCC	AGGCTTACATGACAGGAGGTGGTACCTTTC	2930
Dd	1192	GACGTTCTGCTCC	CCAGTCAGAGGTTCC	AGGCTTACATGACAGGAGGTGGTACCTTTC	1251
Qy	2931	CTGACCAAACTCAG	CAATCAGCTCAGCT	CTCTGTGTAGTCTGACTCTGGCTTACCTATGCT	2990
Dd	1252	CTGACCAAACTCAG	CAATCAGCTCAGCT	CTCTGTGTAGTCTGACTCTGGCTTACCTATGCT	1311
Qy	2991	CCTCTCTCTTCTCT	TATTTCCAGTAAAG	ACCGAGGTCCTGCGCTCTCTCTCTTCCACA	3050
Dd	1312	CCTCTCTCTTCTCT	TATTTCCAGTAAAG	ACCGAGGTCCTGCGCTCTCTCTCTTCCACA	1371

Qy	3051	AGAGTGAGGAGGGCCT	CAGCACACCAACCAT	CATATAGGCCACTTTGAATAGGTCACAAAGG	3110	
Dd	1372	AGAGTGAGGAGGGCCT	CAGCACACCAACCAT	CATATAGGCCACTTTGAATAGGTCACAAAGG	1431	
Qy	3111	CTTTGGCTTCAATTG	AGTAAATACCTTTG	AGTGTATGAGTGAAGCTTTATTTGTTTATC	3170	
Dd	1432	CTTTGGCTTCAATTG	AGTAAATACCTTTG	AGTGTATGAGTGAAGCTTTATTTGTTTATC	1491	
Qy	3171	CATTGAAAGAAATCA	ACTCAAAATTTCTG	TAGATGAGAAAGATGTTGGGAACGAAAAAGG	3230	
Dd	1492	CATTGAAAGAAATCA	ACTCAAAATTTCTG	TAGATGAGAAAGATGTTGGGAACGAAAAAGG	1551	
Qy	3231	CCTAGATAGAGAAAC	AGATCTGCTGAGTA	TAGTACTTAT---GGGGGAGCAGGGGGCG	3286	
Dd	1552	CCTAGATAGAGAAAC	AGATCTGCTGAGTA	TAGTACTTAT---GGGGGAGCAGGGGGCG	1611	
Qy	3287	ATATCCACTGAGTA	CAAGTACTTTGTTG	GGGAGAGAAATCCA	CTGAGTACAAGTACTTGTG	3346
Dd	1612	ATATCCACTGAGTA	CAAGTACTTTGTTG	GGGAGAGAAATCCA	CTGAGTACAAGTACTTGT--	1669
Qy	3347	GCATGGAGATCCACT	GAGTACAAGTACTT	TGTTGGGGGAGGGAATG	GCACAGAGCAAAAGT	3406
Dd	1670	-----	-----	-----GGGGGAAGGAATG	GCACAGAGCAAAAGT	1697
Qy	3407	TGAAGGGA---	AGGAAGATGGAGAG	CGCTCATGTTGGGGGT	TGTAAAGGTCACTCC--TT	3462
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Qy	3523	CAACTATGAACA	TATATCCAGAGG	CGGCGAGACTGTGGG	AGACCTTGGCAATTTAGGGAA	3582
Dd	1818	AA-----	-----	-----CTATGGCAGACT	GTGGAGACCTTGGCAATTTAGGGAA	1855
Qy	3583	GGGGCGCTTTTCA	CGAGAAACTTTAT	GCTCATCTCTTGTG	TACACTCCCACTCCCACTTTG	3642
Dd	1856	GGGGCGCTTTTCA	CGAGAAACTTTAT	GCTCATCTCTTGTG	TACACTCCCACTCCCACTTTG	1915
Qy	3643	ATGAGGTTACGCT	CAGGTTTCTACCG	TTTCTGCTACTG	TGTTGAAAACCTTCAGTAGG	3702
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Dd	1976	ATTCCCAAGAG	CGAGACAGCTCTT	CTGTAAAGGAGGAC	CTTGATTTTCACTGCTCTAG	2035
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Dd	2156	GCTCCCAAGG	ATTAAGATTTCTG	TAGTGTCTTTT	TTTTCAGACATCAGGCT	2215
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## RESULT 5

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US-09-751-797-25
; Sequence 25, Application US/09751797
; Patent No. US20010024652A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: (TIFFS) The Proteins Encoded, and Uses Thereof
; FILE REFERENCES: LUD 5543.2
; CURRENT APPLICATION NUMBER: US/09/751,797
; CURRENT FILING DATE: 2000-12-29
; PRIOR APPLICATION NUMBER: 09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 25
; LENGTH: 4797
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-09-751-797-25
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Query Match 9.2%; Score 686; DB 3; Length 4797;
Best Local Similarity 53.8%; Pred. No. 1.le-144;
Matches 2644; Conservative 0; Mismatches 1875; Indels 393; Gaps 44;
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RESULT 6  
 US-10-627-273-25  
 ; Sequence 25, Application US/10627273  
 ; Publication No. US20040110189A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Dumoutier, Laure  
 ; APPLICANT: Renauld, Jean-Christophe  
 ; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
 ; FILE OF INVENTION: (TIPS) The Proteins Encoded, and Uses Thereof  
 ; FILE REFERENCE: LUD 5543.2  
 ; CURRENT APPLICATION NUMBER: US/10/627,273  
 ; PRIOR FILING DATE: 2003-07-25  
 ; PRIOR APPLICATION NUMBER: US/09/751,797  
 ; PRIOR FILING DATE: 2000-12-29  
 ; PRIOR APPLICATION NUMBER: 09/419,568  
 ; PRIOR FILING DATE: 1999-10-18  
 ; PRIOR APPLICATION NUMBER: US09/178,973  
 ; PRIOR FILING DATE: 1998-10-26  
 ; NUMBER OF SEQ ID NOS: 29  
 ; SEQ ID NO 25  
 ; LENGTH: 4797  
 ; TYPE: DNA  
 ; ORGANISM: Homo sapiens  
 ; FEATURE:  
 US-10-627-273-25

Query Match 9.2%; Score 686; DB 7; Length 4797;  
 Best Local Similarity 53.8%; Pred. No. 1.1e-144;  
 Matches 2644; Conservative 0; Mismatches 1875; Indels 393; Gaps 44;

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 QY 929 TGTCTCTCACTGATAGTTCAGCTTATGACAGGAGTGGTGGCTTCTCTGCGCA 988  
 QY 2938 AACTCAGCAATCAGCTCAGCTCTGTGTAGTGTGACTCTGTGCTACCTATGCTCTCTCT 2997  
 Db |||||  
 QY 989 GGCTCAGCAACAGCTAAGCACATGTGTAACTTCACTCTCAGCTATGCCCACCTACCC 1048  
 QY 2998 CTTCTCTTCTTATTCAGTAAGAACCCGAGGTCTGCGCTCTCTCTCTTCAAGAGTGA 3057  
 Db |||||  
 QY 1049 CTCCTTCCCTCTCTTCCACAGAGACCCCTTACCCCACTCTCTCTCTCTCTCTCTCT 1108  
 QY 3058 GGAGGGCCTCAGCACACACCATATAGSCCATTTGAATAGTGTCAAAAGGCTTTGGC 3117  
 Db |||||  
 QY 1109 TAACTAGCAGGAAGAAGTCTTGGCAGCAGTGTATCAGGAGTCA-----TTTGGG 1161  
 QY 3118 TTCAATTGAGTAACTACTTTGAGTTTGTATGAGTGAAGCTTTATTTGTTTATCCATGGAA 3177  
 Db |||||  
 QY 1162 ATCATAGAGTATTTGCTTTGCTTTGACTGAGTCACTCTTGTAGTTTATAGTGTGAATG 1221  
 QY 3178 AGAAATCAACTCAATTTCTGTAGGATGAGAAAGATGTTGGAAACGAAAAAGGCTTAGAT 3237  
 Db |||||  
 QY 1222 GGTCTGGAACCTTAAGTGTACGAAGCCGATTTGTTGTTCTTGGAAAAAAGCAACTC 1281  
 QY 3238 AGAAAAACAGATCTGCTGAGTAGTACTTATGCGGGGAGCAGGGGCGATATCCACTGA 3297  
 Db |||||  
 QY 1282 A-----GGTTGCGTAAGATGAGAAAGGTGTGG 1309  
 QY 3298 GTACAAAGTACTTGTGGGGAGAGAAATCCACTGAGTACAACTACTTGTGGCATGGAGATC 3357  
 Db |||||  
 QY 1310 GAAAAATCTAGCTGTGGAAATGGATCCATTTAGTCTTGTGAGGGAGGGGATGG 1369  
 QY 3358 CACTGATGACAGTACTTGTGGGGGAGGGAATGGACAGCAAGAAAGTTGAAGGGAGG 3417  
 Db |||||  
 QY 1370 CATGGAGAGAAATTAGAAGAGAAAGTGGGAAATGGGAAGCTTTAA-----1415  
 QY 3418 AAGATGGAGGGCTCATGTTGGGGGTGGAAGGTCACTCTTTTCCATGTGATGGAG 3477  
 Db |||||  
 QY 1416 -----GTGCGTGTGGTGGGTCGCGAGACTGTTGGCTCTGTGATGATCGGGA 1460  
 QY 3478 AGTTAAGAAAAACCA-GTGTGTGAGTTTGTATGTCTTCAGACACCCCCCACTATGAAACAT 3536

Db 1461 AGCCACAAATCGAGGCGTGTGAATTTGATGCCGTGAACATTTGAAATATGAAATAA 1520  
Qy 3537 ATCCACGAGAGCGGCGAGACTGTGGGAGACCTGGGCATTTAGGGAAGCGCG--GGCTTTT 3594  
Db 1521 AGTTTGAGTGGAGTGGGCCAGTAAAGGCCCTAGGACCTTACTGAAGAGGCGCTTAAATTTT 1580  
Qy 3595 CACAGGAAACTTTATGCTCATCTCTGTGCTACACTCCACCTTTGATGAGGTTTCAGC 3654  
Db 1581 CACATGAGATGTTTATGTACATTTCTTGTTCTAAGCATGCAATTTTCTGGAGATACGAT 1640  
Qy 3655 TCAGGTTTTGCTTCT-----ACCGTTCTTGCTACTGTTGGGAAAC 3693  
Db 1641 TGAGGTTTTATTCCTTACAGNAATTTGCATAAATCTACTCCGCTCTTTCCACAAATGCAAC 1700  
Qy 3694 TTCAAGTAGGATTCCTCCAAAGACGAGGACAGCTCTTCTGTGAAGGAGGACCTGGAATTC 3753  
Db 1701 CTCAGTAGGATTTCCCAAGATGAAGAGAGGTCTCTTTGTAAGGGAAGTGACTGGATTCG 1760  
Qy 3754 GTGTCCTAGAGAACGAATAGCTCAGAGAACTTAGGTCACAGTGAATCTAGGTCACAGC 3813  
Db 1761 GCGTCCCAAGGGAATTCAGAGCTCAGGAATCTTAGGTCACGTGTTGAATCTTAGGTCATG 1820  
Qy 3814 GGGCAAAATGACTGAACGCTCTATTCCAGGTGAACGGTCACGCTCAGATATACTG 3873  
Db 1821 TGGGCAAAATTTACTAAGGCTTTAATTCAGGTGAATTTGACTGTACTCTCATCGGTGTG 1880  
Qy 3874 AGGTATTGGGCTCCCGGGAATAGATTCTGTTAGTGA--GTCGCTTTTATTTTGCAGCA 3932  
Db 1881 GAGGTTCAATAAGTTTCAGCAACAATTAAGATAGTTATGCTGTTATGTTTTTATAGCA 1940  
Qy 3933 CATCAGGGTGCAGCACGACGAACTCCAGAGAAATGTCAGAGGCTGAAGGACAGTGAA 3992  
Db 1941 TATTGAAGGTGATGACCTGCATATCCAGAGAAATGTGCAAAAGCTGAAGGACACAGTGA 2000  
Qy 3993 AAAGGTACTATTGGCAAGCCCAATACTAAGCCATTCAGTAG--GAGACGTGGGGATTTC 4050  
Db 2001 AAAGTAGGACTGATACTGTCATGCTCAATGCTAAGTCAATGATGATAGGAGACAAATGTTT 2060  
Qy 4051 TTCTCTGCTCCAGCTCCCTTCTACTTTGTAACATTTTATTTGACTTGTCTACTACTG 4110  
Db 2061 TTCTTTCTCTTTCTTCCCATCACTTTGTGTGATTTTTCATTTGATTTCTCCTACACAG 2120  
Qy 4111 GTCCATTACTCGCTAGTGCACCTGTATCTAGCTGGTCTATAGACTTTTCAATCTGTG 4170  
Db 2121 GCGCATTA-----CTTTGGTGTCTGTGTATGTAGATATATCTATATCTAGATGTCAGTT 2176  
Qy 4171 TCTAAATTT---GTAAGTCACAAATCTCGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTC 4227  
Db 2177 TCCAAATCTTGCAAAATTTAGAAATCTTAGAACTGGTTGGGATCTTAGCTGTCTAGTCAC 2236  
Qy 4228 ATGAGCACTTGTCTGGAGGATGGCTGTGACAGAGTCAATGCTAGAGACAGCATCCCTG 4287  
Db 2237 ATAACCTCAGATTTCTGGGATGGTCAGTGGCAGAGATAGGCTAGATGTCAGGCTCTCTG 2296  
Qy 4288 ATTCACAGCTCTGCAC--TTGCCATGTGGCCATGTGTAAATTAATTTTGGCTTCAATAGTAT 4346  
Db 2297 AATCCCAAGCAGCACATTTTCCCGTGTGTGATACAGATTTAGTTTGTGTACCAATTTCT 2356  
Qy 4347 TTGGGAAA--GCCAGTTCCTCGGACCTTACATATCTGAAGAACCATGCAATTTGAAACTA 4404  
Db 2357 TAGGAAATTTTCAAGTTCTTATGCTCATGTAAATCTGAGAGATGATCTGTTTAAACACA 2416  
Qy 4405 GAAAGCTGGGCA----CAAACCTACTAGAGATGATTTTTTGTAGCTCATTTAAACGGATGCTC 4460  
Db 2417 GAAAAATGCTATGGGCAAAATTTATTTGAAGTCAATTTTTGAAGTCAATTAATGCTTGTCT 2476  
Qy 4461 TGAATGTGGCAAAATCAACCAGAAATTAACAAAGAGCTGGATTTGCAATAGGACA 4520  
Db 2477 TGAACCTTTGGAGAAATAAAGCTCAGAAATGAGAAAGAGCTGGACCTTGCATATAGGGCT 2536  
Qy 4521 AGTATTAGAATCACTGGTATTATTAAGCTATCATCTTAATTAATAATAGGCGCTTATATA 4580

Db 2537 AATTTCTTGA-----GTAATAAACACATTTATTTTGAATATATCAATATCTATCAGATA 2589  
Qy 4581 TATATTTAGATTAACACAGAGTGGATAGCCCTCCCAATTTTACTTTGGCCTGGTTTCAAA 4640  
Db 2590 TTGATTATATAGTTTAAAGCAAGAGCAGACAAC--CGATCTCTTTTATACAGGTTCAAT 2648  
Qy 4641 AGAGTAAAAATATACAGTCATGGATTAAATATAGTGTGCATGAAAGTATGAGATGAAACCC 4700  
Db 2649 AGAGTAAAAATATATAGTAAAGATTTATATAGTAAATGGAAGTCTGAATTTGTAAGCT 2708  
Qy 4701 TTTCTCTACTTTTTCACCTCA-----TTTCTTAGTTTTTTTTTTTCTTTCACACCTGTA 4752  
Db 2709 TTTTTTCTCTCTCTCCCATCAAGACCTTCCATCTAGTTCTTCTTCTTCACTCCCTCA 2768  
Qy 4753 TCAAGCCACTAGTAAGCACCTATCTGCTGTGAGCTATTATGACTTTTACAGCAACAC 4812  
Db 2769 ACAATCCCTTAGGAGCATTTATCCATGTGGGTGGTGTGTACATTTCTATAGTGAATGAT 2828  
Qy 4813 ATTGCTGTGTGGCTCTTTTGGGGAAGGAAACAGATAGCAGGAGGCTCAGGCTAGCAAGT 4872  
Db 2829 ACCATCATGTGGCTATTGTGTGAAGAACA--ACAATGMAAGGCTTAGACTAAACAATA 2886  
Qy 4873 CTGACTTGGCCTTAAAGCCAGAGCATGTGTTGATAGCAGAGAAAGTGAAGGCTCTTCGCAAG 4932  
Db 2887 GTGACTCACCCCAAAACCGGAGGAATGATTAGGAGCAGTGAAGTGAACGCTCTT--GCAAG 2945  
Qy 4933 TGGGTGTGCTTAAGTAATCAGAAACAGGAGGCTCCGGTTGATGCGAATATTCAGTAAGA-- 4991  
Db 2946 CAGGTACAACCTAAATACTCAGAAACATGAAGGCTCCAGTTGATGGAATTTTCAGTAACAA 3005  
Qy 4992 -----TATCTACCCCTTATCTCTTCTTATCGAACCTAAATCGTCTCTTTTTTCTTGTG 5042  
Db 3006 GCTTAACCTTAATTCCTCTTTTCCCTCTGTGACTTTTAAAGAGGCTTTCTTCTCTGAG 3065  
Qy 5043 TGTAGGCTGATAAACACACACTTGT--TTCTTTTTCAGTGTTCATGGCTTTGTAGATTTTAA 5100  
Db 3066 CATCATTTAATGATGTGACTGTGTTCTTCTTTGATAATTTGAAGGCTTTGTAGTTTTAAA 3125  
Qy 5101 GTGCTCTGCCAGTCTTGT--TAGAGGTTTGTGTACCTTCACACCTGGGCTTGGATCTTA 5158  
Db 3126 TTGTGAAGCCAGTCTCTTGTTTATAGAACTATTATCTAGACATGGAAGGCTGGAATGTTA 3185  
Qy 5159 GCATGCCAAAGGACACACTTCTGAATGCTGTGTGAAGAGTTATTATTCATTTACT--- 5215  
Db 3186 GCATGCCACAGACAGCATGCTTTACACATCTTGTCTTAAAGATTTACTGATTTCACTT 3245  
Qy 5216 -----TTGTCTTTGGAAAGGTGAAGCGTGTGTGAAGAAAGAACTCACAGGAGATGTGTTCT 5270  
Db 3246 GCTTGTGTCTTTTAGAAAAGTGAAGTGTGAGAGAGGAATCTCATGGTGA----- 3296  
Qy 5271 CTGTAGGAAAACTTTTTTTTTTCCCTTTAAATGCCCTATAATCCACTTTTCAGTCA---ACTT 5327  
Db 3297 -----TCTGTGTGATTTTCAAGACCTTTTAAATCCATTTTGAAGAAATCAAT 3342  
Qy 5328 TGACTTTTATACCATGCTGCACATGAAGAGTGTTTAGGCCCGCTCTCATGGCTCTGGG 5387  
Db 3343 TCATATTTTGCAATGGGTTGCCATGTGGAAGATGATTTATGCTTTTGTCTGTAGTCTCA 3402  
Qy 5388 AAAAGCAACCAATAGGGAAGGAATGTTTATCTGTAGAAATCTGACCGGAGGAACTGGT 5447  
Db 3403 GAAAGCA--CAGGAGGAGAGCAATGTTGTTTCAAGAAAAGATCAACAGAGGAGGAACCTGT 3461  
Qy 5448 CAGAGCTCCCCGGAAGACCA-----CCACAGGTGTTTAAGTAGG 5485  
Db 3462 CAGAGCTGTCTGAATAATAGGGTGGTTTTGGGAGGCAATTAATTCCTCTCTGTGGGGTAAA 3521  
Qy 5486 AACAGTCCAGGCTGGCTCATGTAATAGAAATGGAACAGAGCGAGGGAAGATAAGCTACAA 5545  
Db 3522 AGCAGAACGCGAGTTGGTAGTAAAT--GCATGACAGACAGTAGGGGAGCAATAACTTTAA 3580  
Qy 5546 AGTTTCATAGGGTC--CGGAGTCTTAAAGATACAAAATAGCTGC--TTGGGCTTCATAACA 5602  
Db 3581 AATTTCTTTATAGTCTTGGAGTCTTTTGAGATAGAAAAGAAATATCTTTTGGCGCTTATGTCA 3640

Qy 5603 AAGGAAGTCTGGGAGGACGCAAGTGAAGGAAATGGAAGGGAAGAAACAGAAATGTAG 5662  
Db 3641 AAGGAAGTATGGAAGG-----TGAAAGGCGGGAAGAAAGCAGGAAAGGAAG 3688  
Qy 5663 AGGACTTGACAGCTACAATCTCTACACAGAGATTTTCTTGGAAACAATCTAGAAGGT 5722  
Db 3689 AACCATGTATATATAGAGACCAATGGTGACAAGGTTTTTCTTTGAAATAATGCAAAATG 3748  
Qy 5723 AGTGGATTAGGTGATTCAGCGGGGACTTGCTTTGCCATTGGAATCTGGGTTTTGTCTCT 5782  
Db 3749 ATAGATTAGAGGAATTTCAAGTGAAGATGCTTTTCACTTGAATTTGGGTTTCTCT--T 3805  
Qy 5783 CCATTGAGTTGAAGCGTCAACCTTTTACCCCTCGAATCGAGAGGAAAGAGGGGTGT 5842  
Db 3806 CGATTAAAGTTGGGATCCTCATCTGCATTTGACT---TGAGAGAGAGAAAGATGAATGT 3861  
Qy 5843 TATGACTCTACCTGGAGTTTACTAGTTTACGGAATGGAACAGACACTCGGACCTCCT 5902  
Db 3862 TAGGACCTATATCTGGTTTTCTATTAACTAAAGCAAGTGGAAAGACTTATTTGGTATTT 3921  
Qy 5903 CTTGACAAAAAATGAAACCTGTTGCTTTGCTGTTGTTGTTCTTTGTTTAAAGAAAGCAC 5962  
Db 3922 TTCCCAAAAAGTGAAACCTTTTCTTTTACTGTTTGTCTTTGTTTGAAGAAAGTGA 3981  
Qy 5963 AGGCAAGCCGACCATCGGTTGAAATGTGGGTCTTTGAGTCAAGGCTTTTGAAGTTGAG 6022  
Db 3982 CCTTAATGTATTGGTGAATACATGTTCAAAAGTCAATTTGAGTAGAGATGTTTAAATCAG 4041  
Qy 6023 CACTCATCAATAGTT-----GATCATGTCAGGTGGAGGGC 6058  
Db 4042 GAGTGTCCAATCAATTTGGCTTCCCTGGACCACTTGAAGAAATTTGCTTGGTACACACAT 4101  
Qy 6059 TACCTGTGAGCGGAGCCCTGCTGGCTTCGCACTTAAACATCTCCAGTCTCAGTATCACT 6118  
Db 4102 AAAATACAGAAACAATAGCTGATGAGCTAAAGAAAGTCCATGCATATAATCTCATACTGTTT 4161  
Qy 6119 TCCTGCTACTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTTTCC----- 6163  
Db 4162 TAAGAAAGTTTATGAATTTCTGTTAGGCTGCAATCAAGGCTGCTGGGCGCATGTGGGC 4221  
Qy 6164 -----AACCCCACTAAATTTAATGACAAAAGACTGTGTAATTTG 6205  
Db 4222 CTGTGGGCTGAGGTTGGACAAGCTCCTTTAAGATATCTGTATAGATAGTTTGGAGC 4281  
Qy 6206 TGGGATACAGTGTGATAATTGA----- 6227  
Db 4282 TGCAAAACAGGCCAAGGCAATATGGGTGGCAGCTCGGGATCCCCAGATCCAGCCTCACT 4341  
Qy 6228 ----TCTATGTGTGCAATGTGCAAGGTTCAATAAGATAGATTAAATAGGCCCATCAACAGC 6283  
Db 4342 TCAGTCTCTTGTCTGTTTAAAGAGGGTGGTCACTCTCTGCCAGCTTTTAAACAGC 4401  
Qy 6284 TTTATGGGTGTGAAATGCAAGTAAATATAGTAGATGCTGTGGTGTCTTTAGTGTGAGAA 6343  
Db 4402 TTTATAGTGTAGGTGCACTGAAATATGATGCTGCTGGTGGCT-CTCAGTCCAGAGA 4460  
Qy 6344 GGCATGATTTTAAAGTCTTGGGCAATCATATATATCTCATGCTATAAATAACATTATGTT 6403  
Db 4461 GCCGTCAATTTAAGCTCTTTGGCAAAATCATACAACTACTAAAGGGATA-----T 4508  
Qy 6404 GATTATTAATCTTTTAGAAGGCGTATCTTGGTTTGGTGTCTCAGCAGCAAAATGTCA 6463  
Db 4509 TACTATGAATGTTTACAAATGTTTAAACTCGTTTCTGTCTCCATCAACCTAATCTTG 4568  
Qy 6464 CCAGCTCTTTTCACTGGTACCACTTTTAGAAAAATGCTGTGCTCAAAATGTTGTTGTA 6523  
Db 4569 CAATTTCT---AATTTGTTTCACTTTAGAAAAACATGTCATAAATGCTCAAAATCTTTTGA 4625  
Qy 6524 TTCTTATTTTCAAGCTTGGAGAGGTGGAGAGATCAGCGGATTTGGGAACTGGAGCTG 6583  
Db 4626 TTCTTATTTTCAAGCTTGGAGAGGTGGAGAGATCAAGCAATTTGGAGAACTGGATTTG 4685

Qy 6584 CTGTTATATGTCTCTGAGAAATGCTTGGTCTGAGCGAGAGAAGCTAGAAAAACGAGAAC 6643  
Db 4686 CTGTTTATGTCTCTGAGAAATGCTTGCATTTGACCAAGCAAGCTGAAAAATGAATAAC 4745  
Qy 6644 TGCTCTCTTCTGCTTCTTAAAAAGAACATAAGATCCCTGAATGGACTTTTTT 6695  
Db 4746 TAACCCCTCTTCTCTGCTGAGAAATGAATTTAGATGCCCAAGCGATTTTTT 4797  
RESULT 7  
US-09-751-797-7  
; Sequence 7, Application US/09751797  
; Patent No. US20010024652A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; (TIFs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/751,797  
; CURRENT FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-751-797-7  
Query Match 8.1%; Score 601.4; DB 3; Length 1119;  
Best Local Similarity 99.8%; Pred. No. 8.4e-126;  
Matches 602; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
Qy 6535 ATAGCTTTGAGAGAGTGGAGAGATCAAGCGATTGGGGAACCTGACCTGCTGTTTATGTC 6594  
Db 510 AAGCTTTGAGAGAGTGGAGAGATCAAGCGATTGGGGAACCTGACCTGCTGTTTATGTC 569  
Qy 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAGCTAGAAAAACGAAAGACTGCTCTTCTCT 6654  
Db 570 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAGCTAGAAAAACGAAAGACTGCTCTTCTCT 629  
Qy 6655 GCCTTCTAAAAAGAACAAATAAGATCCCTGATGAGACTTTTACTTAAAGGAAGTGAAG 6714  
Db 630 GCCTTCTAAAAAGAACAAATAAGATCCCTGATGAGACTTTTACTTAAAGGAAGTGAAG 689  
Qy 6715 GCTAACGTCCTCATCATTTAGAGATTTCACATGAAACCTGGCTCAGTTGAAAAAGAAA 6774  
Db 690 GCTAACGTCCTCATCATTTAGAGATTTCACATGAAACCTGGCTCAGTTGAAAAAGAAA 749  
Qy 6775 TAGTGTCAAGTTGTCTCCATGAGACCAGAGGTAGACTTTGATAACCAACCAAGATTCATTGACA 6834  
Db 750 TAGTGTCAAGTTGTCTCCATGAGACCAGAGGTAGACTTTGATAACCAACCAAGATTCATTGACA 809  
Qy 6835 ATATTTTATGTCACCTGATGATACAAACAGAAAAATTAATGATCTTTAAAAAATTTGTTGAA 6894  
Db 810 ATATTTTATGTCACCTGATGATACAAACAGAAAAATTAATGATCTTTAAAAAATTTGTTGAA 869  
Qy 6895 AGGAGGTTTACCTCTCATCTTCTTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCAA 6954  
Db 870 AGGAGGTTTACCTCTCATCTTCTTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCAA 929  
Qy 6955 TATTTTATATATGTAAGTTTATTTTATTAAGATATACATTTTATTTATGTCAGTTTATTA 7014  
Db 930 TATTTTATATGTAAGTTTATTTTATTAAGATATACATTTTATTTATGTCAGTTTATTA 989  
Qy 7015 ATATGATTTTATATAGAAAAATTAATCTGCTTATGATATTTTAGTATATAAGGCAAAATA 7074  
Db 990 ATATGATTTTATTTATAGAAAAATTAATCTGCTTATGATATTTTAGTATATAAGGCAAAATA 1049

QY 7075 TTTATGACAACTATGGAACAGATATCTTAGGCTTTTAATAAACAACATGATATCAT 7134  
Db 1050 TTTATGACAACTATGGAACAGATATCTTAGGCTTTTAATAAACAACATGATATCAT 1109  
QY 7135 AAA 7137  
Db 1110 AAA 1112

RESULT 8  
US-10-627-273-7  
; Sequence 7, Application US/10627273  
; Publication No. US20040110189A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/10/627,273  
; PRIOR FILING DATE: 2003-07-25  
; PRIOR FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-10-627-273-7

Query Match 8.1%; Score 601.4; DB 7; Length 1119;  
Best Local Similarity 99.8%; Pred. No. 8.4e-126;  
Matches 602; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 6535 ATAGCTTGGAGAGAGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 6594  
Db 510 AAGCTTGGAGAGAGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 569  
QY 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAAAACGAAGAACTGCTCCTTCCT 6654  
Db 570 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAAAACGAAGAACTGCTCCTTCCT 629  
QY 6655 GCCTTCTAAAAAGAACAAATAAGATCCCTGAATGGACCTTTTACTAAAGGAAAGTGAGAA 6714  
Db 630 GCCTTCTAAAAAGAACAAATAAGATCCCTGAATGGACCTTTTACTAAAGGAAAGTGAGAA 689  
QY 6715 GCTAACGTCCTCATCTATTAGAGATTTTCACATGAAACCTGGCTCAGTTGAAAAAGAAAA 6774  
Db 690 GCTAACGTCCTCATCTATTAGAGATTTTCACATGAAACCTGGCTCAGTTGAAAAAGAAAA 749  
QY 6775 TAGTGTCAAGTTGTCCATGAGACAGAGGTAGACTTGTATAACCAAGAGATTCATTGACA 6834  
Db 750 TAGTGTCAAGTTGTCCATGAGACAGAGGTAGACTTGTATAACCAAGAGATTCATTGACA 809  
QY 6835 ATATTTTATGTCACTGATGATACAAACAGAAAAATAATGTACTTTTAAAAAATTTGTTGAA 6894  
Db 810 ATATTTTATGTCACTGATGATACAAACAGAAAAATAATGTACTTTTAAAAAATTTGTTGAA 869  
QY 6895 AGGAGTTACCTCTCATCTTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCAA 6954  
Db 870 AGGAGTTACCTCTCATCTTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCAA 929  
QY 6955 TATTTTATATGTAAGTTTATTTATTAAGTATACATATTTTATTTATGTGCTAGTTTATTA 7014  
Db 930 TATTTTATATGTAAGTTTATTTATTAAGTATACATATTTTATTTATGTGCTAGTTTATTA 989

QY 7015 ATATGATTTTATTTATAGAAACATTTATCTGCTATTTGATATTTTAGTATAAGGCAAAATAA 7074  
Db 990 ATATGATTTTATTTATAGAAACATTTATCTGCTATTTGATATTTTAGTATAAGGCAAAATAA 1049  
QY 7075 TTTATGACAACTATGGAACAGATATCTTAGGCTTTTAATAAACAACATGATATCAT 7134  
Db 1050 TTTATGACAACTATGGAACAGATATCTTAGGCTTTTAATAAACAACATGATATCAT 1109  
QY 7135 AAA 7137  
Db 1110 AAA 1112

RESULT 9  
US-10-084-298-3  
; Sequence 3, Application US/10084298  
; Publication No. US20030099649A1  
; GENERAL INFORMATION:  
; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: Xuan, DeJun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory  
; FILE REFERENCE: GI5358 CIP  
; CURRENT APPLICATION NUMBER: US/10/084,298  
; CURRENT FILING DATE: 2002-09-10  
; PRIOR APPLICATION NUMBER: 60/270,823  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR APPLICATION NUMBER: 60/281,353  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR APPLICATION NUMBER: 60/131,473  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR APPLICATION NUMBER: 09/561,811  
; PRIOR FILING DATE: 2000-04-28  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 3  
; LENGTH: 1166  
; TYPE: DNA  
; ORGANISM: Murine  
US-10-084-298-3

Query Match 8.0%; Score 598.2; DB 5; Length 1166;  
Best Local Similarity 99.5%; Pred. No. 4.6e-125;  
Matches 600; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 6535 ATAGCTTGGAGAGAGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 6594  
Db 533 AAGCTTGGAGAGAGTGGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 592  
QY 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAAAACGAAGAACTGCTCCTTCCT 6654  
Db 593 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAAAACGAAGAACTGCTCCTTCCT 652  
QY 6655 GCCTTCTAAAAAGAACAAATAAGATCCCTGAATGGACCTTTTACTAAAGGAAAGTGAGAA 6714  
Db 653 GCCTTCTAAAAAGAACAAATAAGATCCCTGAATGGACCTTTTACTAAAGGAAAGTGAGAA 712  
QY 6715 GCTAACCTCCATCATCTATTAGAGATTTTCATGAAACCTGGCTCAGTTGAAAAAGAAAA 6774  
Db 713 GCTAACCTCCATCATCTATTAGAGATTTTCATGAAACCTGGCTCAGTTGAAAAAGAAAA 772  
QY 6775 TAGTGTCAAGTTGTCCATGAGACAGAGGTAGACTTGTATAACCAAGAGATTCATTGACA 6834  
Db 773 TAGTGTCAAGTTGTCCATGAGACAGAGGTAGACTTGTATAACCAAGAGATTCATTGACA 832  
QY 6835 ATATTTTATGTCACTGATGATACAAACAGAAAAATAATGTACTTTTAAAAAATTTGTTGAA 6894  
Db 833 ATATTTTATGTCACTGATGATACAAACAGAAAAATAATGTACTTTTAAAAAATTTGTTGAA 892  
QY 6895 AGGAGTTACCTCTCATCTTTTAGAAAAAGCTTATGTAACCTTCATTTCCATATCCAA 6954

Db 893 AGGAGGTTACCTCTCATTCTCTTTAGAAAAAAGCTTATGTAACCTTCAATTTCCATAACCA 952  
Qy 6955 TATTATATATAGTAAGTATTATTATTATAAGTATACATTTTATTATGTGAGTTTATTA 7014  
Db 953 TATTATATATAGTAAGTATTATTATTATAAGTATACATTTTATTATGTGAGTTTATTA 1012  
Qy 7015 ATATGAGTTTATTATATAGAACATTTCTGCTATGTGATATTTAGTATATAAGGCAAAATA 7074  
Db 1013 ATATGAGTTTATTATATAGAACATTTCTGCTATGTGATATTTAGTATATAAGGCAAAATA 1072  
Qy 7075 TTTATGACATACTATGGAACAAGATATCTTAGGCTTTTAAATAACACATGGATATCAT 7134  
Db 1073 TTTATGACATACTATGGAACAAGATATCTTAGGCTTTTAAATAACACATGGATATCAT 1132  
Qy 7135 AAA 7137  
Db 1133 AAA 1135

## RESULT 10

US-10-256-977-3  
; Sequence 3, Application US/10256977  
; Publication No. US20030157106A1  
; GENERAL INFORMATION:  
; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: Xuan, Dejun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory  
; FILE OF INVENTION: Disorders  
; FILE REFERENCE: GI5358 CIP  
; CURRENT APPLICATION NUMBER: US/10/256,977  
; CURRENT FILING DATE: 2002-09-27  
; PRIOR APPLICATION NUMBER: US/10/084,298  
; PRIOR FILING DATE: 2002-09-10  
; PRIOR APPLICATION NUMBER: 60/270,823  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR APPLICATION NUMBER: 60/281,353  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR APPLICATION NUMBER: 60/131,473  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR APPLICATION NUMBER: 09/561,811  
; PRIOR FILING DATE: 2000-04-28  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 3  
; LENGTH: 1166  
; TYPE: DNA  
; ORGANISM: Murine

## US-10-256-977-3

Query Match 8.0%; Score 598.2; DB 6; Length 1166;  
Best Local Similarity 99.5%; Pred. No. 4.6e-125;  
Matches 600; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 6535 ATAGCTTGGAGAGAGTGAGAGATCAAGGCGATTGGGGAACCTGACCTGCTGTTTATGTC 6594  
Db 533 AAGCTTGGAGAGAGTGAGAGATCAAGGCGATTGGGGAACCTGACCTGCTGTTTATGTC 592  
Qy 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAACCGAAGAACTGCTCCTTCT 6654  
Db 593 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAACCGAAGAACTGCTCCTTCT 652  
Qy 6655 GCCTTCTAAAGAAACAATAAGATCCCTGATGACATTTTACTTAAAGAAAGTGAGAA 6714  
Db 653 GCCTTCTAAAGAAACAATAAGATCCCTGATGACATTTTACTTAAAGAAAGTGAGAA 712  
Qy 6715 GCTAACCTCCATCATCATATAGAGATTTCATGAAACCTGGCTCAGTTCGAAAAGAA 6774  
Db 713 GCTAACCTCCATCATCATATAGAGATTTCATGAAACCTGGCTCAGTTCGAAAAGAA 772

Qy 6775 TAGTGCAAGTTGCTCATGAGACCAGAGGTAGAGCTTGATTAACCAAGATTTCAATTGACA 6834  
Db 773 TAGTGCAAGTTGCTCATGAGACCAGAGGTAGAGCTTGATTAACCAAGATTTCAATTGACA 832  
Qy 6835 ATATTTTATTGTGCACGTGATACAAACAGAAAAAATATGTACTTTTAAAAAATTTGTTTGA 6894  
Db 833 ATATTTTATTGTGCACGTGATACAAACAGAAAAAATATGTACTTTTAAAAAATTTGTTTGA 892  
Qy 6895 AGGAGGTTACCTCTCATTCTCTTTAGAAAAAAGCTTATGTAACCTTCAATTTCCATAACCA 6954  
Db 893 AGGAGGTTACCTCTCATTCTCTTTAGAAAAAAGCTTATGTAACCTTCAATTTCCATAACCA 952  
Qy 6955 TATTATATATAGTAAGTATTATTATTATAAGTATACATTTTATTATGTGAGTTTATTA 7014  
Db 953 TATTATATATAGTAAGTATTATTATTATAAGTATACATTTTATTATGTGAGTTTATTA 1012  
Qy 7015 ATATGAGTTTATTATATAGAACATTTCTGCTATGTGATATTTAGTATATAAGGCAAAATA 7074  
Db 1013 ATATGAGTTTATTATATAGAACATTTCTGCTATGTGATATTTAGTATATAAGGCAAAATA 1072  
Qy 7075 TTTATGACATACTATGGAACAAGATATCTTAGGCTTTTAAATAACACATGGATATCAT 7134  
Db 1073 TTTATGACATACTATGGAACAAGATATCTTAGGCTTTTAAATAACACATGGATATCAT 1132  
Qy 7135 AAA 7137  
Db 1133 AAA 1135

## RESULT 11

US-10-873-972-3  
; Sequence 3, Application US/10873972  
; Publication No. US20050042220A1  
; GENERAL INFORMATION:  
; APPLICANT: Li, Jing  
; APPLICANT: Tan, Xiang-Yang  
; APPLICANT: Tomkinson, Kathleen N.  
; APPLICANT: Pittman, Debra D.  
; APPLICANT: Veldman, Geertruida M.  
; APPLICANT: Fouser, Lynette  
; TITLE OF INVENTION: Antibodies Against Interleukin-22 and Uses Therefor  
; FILE REFERENCE: AM01524  
; CURRENT APPLICATION NUMBER: US/10/873,972  
; CURRENT FILING DATE: 2004-06-22  
; PRIOR APPLICATION NUMBER: US 60/480,652  
; PRIOR FILING DATE: 2003-06-23  
; PRIOR APPLICATION NUMBER: US 10/084,298  
; PRIOR FILING DATE: 2002-02-25  
; PRIOR APPLICATION NUMBER: US 60/270,823  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR APPLICATION NUMBER: US 60/281,353  
; PRIOR FILING DATE: 2001-04-03  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: PatentIn version 3.1

; SEQ ID NO 3  
; LENGTH: 1166  
; TYPE: DNA  
; ORGANISM: Mus musculus

## US-10-873-972-3

Query Match 8.0%; Score 598.2; DB 8; Length 1166;  
Best Local Similarity 99.5%; Pred. No. 4.6e-125;  
Matches 600; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 6535 ATAGCTTGGAGAGAGTGAGAGATCAAGGCGATTGGGGAACCTGACCTGCTGTTTATGTC 6594  
Db 533 AAGCTTGGAGAGAGTGAGAGATCAAGGCGATTGGGGAACCTGACCTGCTGTTTATGTC 592  
Qy 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAACCGAAGAACTGCTCCTTCT 6654  
Db 593 TCTGAGAAATGCTTGGCTCTGAGCGAGAAGAGCTAGAAACCGAAGAACTGCTCCTTCT 652  
Qy 6655 GCCTTCTAAAGAAACAATAAGATCCCTGATGACATTTTACTTAAAGAAAGTGAGAA 6714

Db 653 GCCTTCTAAAAGAACAAATAGATCCCTGATGACCTTTTACTAAAGGAAAGTGAGAA 712  
Qy 6715 GCTAACGTCATCATATAGAGATTTTACATGAAACCTGGCTCAGTTGAAAAGAAAA 6774  
Db 713 GCTAACGTCATCATATAGAGATTTTACATGAAACCTGGCTCAGTTGAAAAGAAAA 772  
Qy 6775 TAGTGCAAGTTGTCATGAGACGAGAGGAGAGCTTGCATTAACCAAAAGATTCATTGACA 6834  
Db 773 TAGTGCAAGTTGTCATGAGACGAGAGGAGAGCTTGCATTAACCAAAAGATTCATTGACA 832  
Qy 6835 ATATTTTATTTGCTACTGATGATACACAGAAAAATAATATGTAATTTTAAAAAATGTTTGAA 6894  
Db 833 ATATTTTATTTGCTACTGATGATACACAGAAAAATAATATGTAATTTTAAAAAATGTTTGAA 892  
Qy 6895 AGGAGGTACCTCTCATCTCTTTAGAAAAAAGCTTATGTAATTTTCCATATCCAA 6954  
Db 893 AGGAGGTACCTCTCATCTCTTTAGAAAAAAGCTTATGTAATTTTCCATATCCAA 952  
Qy 6955 TATTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 7014  
Db 953 TATTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 1012  
Qy 7015 ATATGGATTTTATTAAGAACATTTATCTGCTATTGATATTTTAGTATAAGGCCAAATAATA 7074  
Db 1013 ATATGGATTTTATTAAGAACATTTATCTGCTATTGATATTTTAGTATAAGGCCAAATAATA 1072  
Qy 7075 TTTATGACATAACTATGGAACAAAGATATCTTAGGCTTTAATAAACACATGGATATCAT 7134  
Db 1073 TTTATGACATAACTATGGAACAAAGATATCTTAGGCTTTAATAAACACATGGATATCAT 1132  
Qy 7135 AAA 7137  
Db 1133 AAA 1135

RESULT 12  
US-11-157-387-3  
; Sequence 3, Application US/11157387  
; Publication No. US20050238648A1  
; GENERAL INFORMATION:  
; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: Xuan, Dejun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory Disorders  
; FILE REFERENCE: G15358 CIP  
; CURRENT FILING DATE: 2005-06-20  
; PRIOR APPLICATION NUMBER: US/11/157,387  
; PRIOR FILING DATE: 2002-09-10  
; PRIOR APPLICATION NUMBER: 60/270,823  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR APPLICATION NUMBER: 60/281,353  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR APPLICATION NUMBER: 60/131,473  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR APPLICATION NUMBER: 09/561,811  
; PRIOR FILING DATE: 2000-04-28  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 3  
; LENGTH: 1166  
; TYPE: DNA  
; ORGANISM: Murine  
US-11-157-387-3

Query Match 8.0%; Score 598.2; DB 10; Length 1166;  
Best Local Similarity 99.5%; Pred. No. 4.6e-125;  
Matches 600; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 6535 ATAGCTTGGAGAGAGTGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 6594  
Db 533 AAAGCTTGGAGAGAGTGAGAGATCAAGGCGATTGGGGAACCTGGACCTGCTGTTTATGTC 592  
Qy 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGAGCTAGAAAACGAAGAACTGCTCTTCCCT 6654  
Db 593 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGAGCTAGAAAACGAAGAACTGCTCTTCCCT 652  
Qy 6655 GCCTTCTTAAAAAGAACAAATAAGATCCCTGAATGACCTTTTACTTAAAGGAAAGTGAGAA 6714  
Db 653 GCCTTCTTAAAAAGAACAAATAAGATCCCTGAATGACCTTTTACTTAAAGGAAAGTGAGAA 712  
Qy 6715 GCTAAAGCTCCATCATCATTTAGAAAGATTTTCAATGAAAAACCTGGCTCAGTTGAAAAAGAAAA 6774  
Db 713 GCTAAAGCTCCATCATTTAGAAAGATTTTCAATGAAAAACCTGGCTCAGTTGAAAAAGAAAA 772  
Qy 6775 TAGTGCAAGTTGTCATGAGACGAGAGGAGAGCTTGCATTAACCAAAAGATTCATTGACA 6834  
Db 773 TAGTGCAAGTTGTCATGAGACGAGAGGAGAGCTTGCATTAACCAAAAGATTCATTGACA 832  
Qy 6835 ATATTTTATTTGCTACTGATGATACACAGAAAAATAATGTAATTTTAAAAAATGTTTGAA 6894  
Db 833 ATATTTTATTTGCTACTGATGATACACAGAAAAATAATGTAATTTTAAAAAATGTTTGAA 892  
Qy 6895 AGGAGGTACCTCTCATCTCTTTAGAAAAAAGCTTATGTAATTTTCCATATCCAA 6954  
Db 893 AGGAGGTACCTCTCATCTCTTTAGAAAAAAGCTTATGTAATTTTCCATATCCAA 952  
Qy 6955 TATTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 7014  
Db 953 TATTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 1012  
Qy 7015 ATATGGATTTTATTAAGAACATTTATCTGCTATTGATATTTTAGTATAAGGCCAAATAATA 7074  
Db 1013 ATATGGATTTTATTAAGAACATTTATCTGCTATTGATATTTTAGTATAAGGCCAAATAATA 1072  
Qy 7075 TTTATGACATAACTATGGAACAAAGATATCTTAGGCTTTAATAAACACATGGATATCAT 7134  
Db 1073 TTTATGACATAACTATGGAACAAAGATATCTTAGGCTTTAATAAACACATGGATATCAT 1132  
Qy 7135 AAA 7137  
Db 1133 AAA 1135

RESULT 13  
US-09-751-797-9  
; Sequence 9, Application US/09751797  
; Patent No. US20010024652A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors and Uses Thereof  
; TITLE OF INVENTION: (TIFF) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/751,797  
; CURRENT FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-751-797-9

Query Match 7.5%; Score 555.2; DB 3; Length 1111;  
Best Local Similarity 96.0%; Pred. No. 2.8e-115;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;



; NAME/KEY: CDS  
; LOCATION: (50)...(589)  
US-10-090-365-40

Query Match		6.7%;	Score 499;	DB 5;	Length 1050;	
Best Local Similarity		96.1%;	Pred. No. 1.7e-102;			
Matches 522;		Conservative 0;	Mismatches 20;	Indels 1;	Gaps 1;	
Qy	6535	ATAGCTTGGAGAGAGTGGAGAGATCAAGGGGATTTGGGGAACTGGACCTGCTGTTTATGTC	6594			
Db	508	AAAGCTTGGAGAGAGCGAGAGATCAAAAGCGATCGGGGAACTGGACCTGCTGTTTATGTC	567			
Qy	6595	TCTCAGAAATGCTTGGCTCTGAGCGAGAAGCTAGAAAACGAAGAACTGCTCCTTCCT	6654			
Db	568	TCTCAGAAATGCTTGGCTCTGAGCGAGAAGCTAGAAAACGAAGAACTGCTCCTTCCT	627			
Qy	6655	GCCTTCTAAAGAAACAATAAGATCCCTGAAATGGAATTTTTTACTAAAGGAAAGTGAGAA	6714			
Db	628	GCCTTCTAAAGAAACAATAAGATCCCTGAAATGGAATTTTTTACTAAAGGAAAGTGAGAA	687			
Qy	6715	GCTAACGTCCATCATCATTTAGAAAGATTTACATGAAACCTGGCTCAGTTGAAAAGAAAA	6774			
Db	688	GCTAACGTCCACCATCATTTAGAAAGATTTACATGAAACCTGGCTCAGTTGAAAAGAAAA	747			
Qy	6775	TAGTGTCAGGTGTGCTCATGAGACGAGAGGTAGACTTGTATACCAAAAGATTTCATTGACA	6834			
Db	748	TAGTGTCAGGTGTGCTCATGAGACGAGAGGTAGACTTGTATACCAAAAGATTTCATTGACA	807			
Qy	6835	ATATTTTATTTGCTCACTGATGATACACAGAAAAATAATGTACTTTTAAAAAATTTGTTGAA	6894			
Db	808	ATATTTTATTTGCTCACTGATGATACACAGAAAAATAATGTACTTTTAAAAAATTTGTTGAA	867			
Qy	6895	AGGAGGTTACCTCTCATTTCTTTAGAAAAAAGCTTATGTAACCTTCCATATCCAA	6954			
Db	868	AGGAGGTTACCTCTCATTTCTTTAGAAAAAAGCTTATGTAACCTTCCATATCCAA	927			
Qy	6955	TATTTTATATATGTAAGTTTATTTATTTATATATAGTATACATTTTATTTGTCAGTTTATTA	7014			
Db	928	TATTTTATATATGTAAGTTTATTTATTTATATATAGTATACATTTTATTTGTCAGTTTATTA	987			
Qy	7015	ATATGGATTTTATTTATAGAAACATTATCTGCTATTGTAATTT-AGTATAAGGCAAAATAAT	7073			
Db	988	ATATGGATTTTATTTATAGAAAAATTAICTGATGTTGATATTTTGAGTATAAGCAAAATAAT				
Qy	7074	ATT 7076				
Db	1048	ATT 1050				

Search completed: December 21, 2005, 21:08:10  
Job time : 3788.38 secs



GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 519.107 Seconds  
(without alignments)  
7442.822 Million cell updates/sec

Title: US-09-751-797-8

Perfect score: 7445

Sequence: 1 gctatcacctgcttaagat.....gattaattaattgtgat 7445

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 4168288 segs, 259477437 residues

Total number of hits satisfying chosen parameters: 8336576

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published Applications\_NA\_New.\*  
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2: /cgn2\_6/ptodata/2/pubpna/US06\_NEW\_PUB.seq.\*  
3: /cgn2\_6/ptodata/2/pubpna/US07\_NEW\_PUB.seq.\*  
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5: /cgn2\_6/ptodata/2/pubpna/US09\_NEW\_PUB.seq.\*  
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7: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq.\*  
8: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq2.\*  
9: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq3.\*  
10: /cgn2\_6/ptodata/2/pubpna/US60\_NEW\_PUB.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	7441.8	100.0	7445	7 US-11-177-987-8	Sequence 8, Appli
2	4245.2	57.0	5935	7 US-11-177-987-42	Sequence 42, Appl
3	686	9.2	4797	7 US-11-177-987-26	Sequence 26, Appl
4	601.4	8.1	1119	7 US-11-177-987-7	Sequence 7, Appli
5	555.2	7.5	1111	7 US-11-177-987-9	Sequence 9, Appli
6	214.2	2.9	1152	7 US-11-102-240-153	Sequence 153, App
7	126	1.7	690	7 US-11-177-987-25	Sequence 25, Appl
8	122	1.6	418	7 US-11-177-987-18	Sequence 18, Appl
9	70	0.9	17395	6 US-10-995-561-13277	Sequence 13277, A
10	62.6	0.8	119036	6 US-10-995-561-13314	Sequence 13314, A
11	61.6	0.8	11462	7 US-11-140-417-22	Sequence 22, Appl
12	59.2	0.8	171936	6 US-10-933-025-24	Sequence 24, Appl
13	58.4	0.8	171486	7 US-11-121-086-105	Sequence 105, App
14	57.4	0.8	20317	6 US-10-995-561-13460	Sequence 13460, A
15	57.2	0.8	173602	7 US-11-121-086-25	Sequence 25, Appl
16	57.2	0.8	184868	7 US-11-121-086-88	Sequence 88, Appl
17	56.6	0.8	201990	6 US-10-995-561-13303	Sequence 13303, A
18	56.4	0.8	196	6 US-10-502-972-25	Sequence 25, Appl
19	56	0.8	49979	6 US-10-995-561-13443	Sequence 13443, A
20	56	0.8	171486	7 US-11-121-086-105	Sequence 105, App
21	54.8	0.7	431	7 US-11-108-172-584	Sequence 584, App
22	54.8	0.7	1652	7 US-11-159-516A-28	Sequence 28, Appl
23	54	0.7	139054	7 US-11-121-086-96	Sequence 96, Appl

## ALIGNMENTS

## RESULT 1

US-11-177-987-8

; Sequence 8, Application US/11177987

; Publication No. US20050271619A1

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5664

; CURRENT FILING DATE: 2005-07-07

; PRIOR APPLICATION NUMBER: US/11/177,987

; PRIOR FILING DATE: 2000-07-27

; PRIOR APPLICATION NUMBER: US09/419,568

; PRIOR FILING DATE: 1999-10-18

; PRIOR APPLICATION NUMBER: US09/354,243

; PRIOR FILING DATE: 1999-07-16

; PRIOR APPLICATION NUMBER: US09/178,973

; PRIOR FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 43

; SEQ ID NO 8

; LENGTH: 7445

; TYPE: DNA

; ORGANISM: Mus musculus

; FEATURE:

US-11-177-987-8

Query Match 100.0%; Score 7441.8; DB 7; Length 7445;

Best Local Similarity 100.0%; Pred. No. 0;

Matches 7443; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy	1	GTCTATCACCCTGCTTAAGATCTTCTTAATTTATAAAAAAACAATTTCTTAAATGAAAA	60
Db	1	GTCTATCACCCTGCTTAAGATCTTCTTAATTTATAAAAAAACAATTTCTTAAATGAAAA	60
Qy	61	GCACACAGCAGCAGTATTTATAGCATGGTCTTGACCATGCAGGTACAGAGTGAATGG	120
Db	61	GCAACAGAGCAGCAGTATTTATAGCATGGTCTTGACCATGCAGGTACAGAGTGAATGG	120
Qy	121	TAAGAGCGCGCTATTATCAGCATTAACCAACATGTTTCTTCTTCTGCAAGCAAACT	180
Db	121	TAGAGCGCGCTATTATCAGCATTAACCAACATGTTTCTTCTTCTGCAAGCAAACT	180
Qy	181	TGAAATCTATGCTTTTAAACAATCTTCAAGCCTCTAATATATAGTGTCTAACGACTGAGTCCG	240

Db 181 TGAATCTATGTCCTTAAACAATCTTCAAGCCTCTAATATAGTGTCTAAACGACTGGAGTCG 240  
 Qy 241 CTGCTGTCCAAACAGAGCTCTTGACACGCTCTCTCTGTTGGCAATTTTATGTTCTTTGA 300  
 Db 241 CTGCTGTCCAAACAGAGCTCTTGACACGCTCTCTCTGTTGGCAATTTTATGTTCTTTGA 300  
 Qy 301 TCGACTCCCAACCTCTCACCTTCGGCTCTGATGGCCACCTTTCAACTTTTCTGCAATTTA 360  
 Db 301 TCGACTCCCAACCTCTCACCTTCGGCTCTGATGGCCACCTTTCAACTTTTCTGCAATTTA 360  
 Qy 361 TGAATCCATGTTTAAATCTTTTAAATAATATTCACAATACTGTTTGTGCAAGTC 420  
 Db 361 TGAATCCATGTTTAAATCTTTTAAATAATATTCACAATACTGTTTGTGCAAGTC 420  
 Qy 421 TGTTCACCCACATGATGTCGTGACCAAGTCTGCTGGTGGTCTGTTGGGGGCAAGGA 480  
 Db 421 TGTTCACCCACATGATGTCGTGACCAAGTCTGCTGGTGGTCTGTTGGGGGCAAGGA 480  
 Qy 481 GCAGGAGAGGTGCCCTGGCACCGGAGTCAAGGATGTTGTAGCTTCCACCTAGAAATGAGATGCT 540  
 Db 481 GCAGGAGAGGTGCCCTGGCACCGGAGTCAAGGATGTTGTAGCTTCCACCTAGAAATGAGATGCT 540  
 Qy 541 GGGAGTTAGACCCAGGTCCTCCAGAAGTGCAGCAAAATGCTCTTAAACACACGCAAGCAAT 600  
 Db 541 GGGAGTTAGACCCAGGTCCTCCAGAAGTGCAGCAAAATGCTCTTAAACACACGCAAGCAAT 600  
 Qy 601 TCTCTCTCAGCCCAACATGAGTGTCTTTAGATTTCCACCTAGAAATGAGATGCTGAGTGC 660  
 Db 601 TCTCTCTCAGCCCAACATGAGTGTCTTTAGATTTCCACCTAGAAATGAGATGCTGAGTGC 660  
 Qy 661 TTCACTCACTGCCACCTCCCTTTCGATCTTTCTGCAAGTGCAGCAAAATGCTCTTAAACACACGCAAGCAAT 720  
 Db 661 TTCACTCACTGCCACCTCCCTTTCGATCTTTCTGCAAGTGCAGCAAAATGCTCTTAAACACACGCAAGCAAT 720  
 Qy 721 CCCACACTGTCTTCGCTCTCAAGTCTGCACTCTCAACAGGTCAAGATTTCTCCAGTGT 780  
 Db 721 CCCACACTGTCTTCGCTCTCAAGTCTGCACTCTCAACAGGTCAAGATTTCTCCAGTGT 780  
 Qy 781 CCCTCTAACACTTTCGAGTGCCTCTTAACACTTTCTCAAGTGTCCCTCTAACACTTTT 840  
 Db 781 CCCTCTAACACTTTCGAGTGCCTCTTAACACTTTCTCAAGTGTCCCTCTAACACTTTT 840  
 Qy 841 CTCAGTGTCCCTCTAACACTTTTGAATCTCAATAGCTGAGGGAGAAAGATCTCACACA 900  
 Db 841 CTCAGTGTCCCTCTAACACTTTTGAATCTCAATAGCTGAGGGAGAAAGATCTCACACA 900  
 Qy 901 GTGATTTTCATGACTTCGCGTTCTAGTCTAGATGTAGGCAATTTGCGTGTCAAGTCTAGGGT 960  
 Db 901 GTGATTTTCATGACTTCGCGTTCTAGTCTAGATGTAGGCAATTTGCGTGTCAAGTCTAGGGT 960  
 Qy 961 AGGCTGTGCTCCGCTGCTTAGGAAGACTTTTCTAGTCTAGTGTGCTGAGTGTCTATCTG 1020  
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 Qy 1021 GGAATCAGTGTACATACAAATCCAGATTTTCTGTAATTTCTCTTCAACT 1080  
 Db 1021 GGAATCAGTGTACATACAAATCCAGATTTTCTGTAATTTCTCTTCAACT 1080  
 Qy 1081 ATCCATCTATAGTATGTTAGTGTCTCAATTTTAAATAATATTTTGAAGCTTATGC 1140  
 Db 1081 ATCCATCTATAGTATGTTAGTGTCTCAATTTTAAATAATATTTTGAAGCTTATGC 1140  
 Qy 1141 TTGCACAAAGTAAATGTCCAGAAATTAGCAATGTATAGTATTTTATTTTAAAAAA 1200  
 Db 1141 TTGCACAAAGTAAATGTCCAGAAATTAGCAATGTATAGTATTTTATTTTAAAAAA 1200  
 Qy 1201 TCTATGCTTAAATGTCTATAGTGTCTCACTACCGATATTTCCAACTTAACTTGACC 1260  
 Db 1201 TCTATGCTTAAATGTCTATAGTGTCTCACTACCGATATTTCCAACTTAACTTGACC 1260  
 Qy 1261 TTGGCTATGATTTCAACCTTTGTATTTGCACTCTACCAATACAGTCTCTGAAACCAAGACAT 1320  
 Db 1261 TTGGCTATGATTTCAACCTTTGTATTTGCACTCTACCAATACAGTCTCTGAAACCAAGACAT 1320

Qy 1321 TCTGTGGCAATGGAGCTGTGAAGAAAGCCAACTTCTTATTTAAAAAAGGAGGCTT 1380  
 Db 1321 TCTGTGGCAATGGAGCTGTGAAGAAAGCCAACTTCTTATTTAAAAAAGGAGGCTT 1380  
 Qy 1381 GTTATAGTTTGGATTCCATATATAAATAAATAGAGATATATTTATTTTAAAAATGA 1440  
 Db 1381 GTTATAGTTTGGATTCCATATATAAATAAATAGAGATATATTTATTTTAAAAATGA 1440  
 Qy 1441 AATAATCTCAAGTTTTCATTATGGCTTATTTCAAGCAAGATATAGGACAGGCTCT 1500  
 Db 1441 AATAATCTCAAGTTTTCATTATGGCTTATTTCAAGCAAGATATAGGACAGGCTCT 1500  
 Qy 1501 TTTATTTCTGCTCACTTCTTAAAGAGATAAGAAATCTATGAAGTTGGTGGGAAATAGAGTCC 1560  
 Db 1501 TTTATTTCTGCTCACTTCTTAAAGAGATAAGAAATCTATGAAGTTGGTGGGAAATAGAGTCC 1560  
 Qy 1561 GTGACCAAAAACGCTGACTCAATAGCTACGGGAGATCAAGGCTGCTCTACTCAATCAGAA 1620  
 Db 1561 GTGACCAAAAACGCTGACTCAATAGCTACGGGAGATCAAGGCTGCTCTACTCAATCAGAA 1620  
 Qy 1621 TCTACTACGGCAAAAGCCATGGCTTTCTTTGAAAACCGTGTTTAGAAAGATTTCTGGGATTT 1680  
 Db 1621 TCTACTACGGCAAAAGCCATGGCTTTCTTTGAAAACCGTGTTTAGAAAGATTTCTGGGATTT 1680  
 Qy 1681 GTGTGCAAAAAGCACCTTGTGTGGCCCTCACCGTGACGTTTGTAGGGAAGACTTCCCATCTCT 1740  
 Db 1681 GTGTGCAAAAAGCACCTTGTGTGGCCCTCACCGTGACGTTTGTAGGGAAGACTTCCCATCTCT 1740  
 Qy 1741 CAAGGTGGGAAGGCTTGGAGGTGGTGTCTTGTGGCCCTCTATGGTGGTTAGGTAATTTCTC 1800  
 Db 1741 CAAGGTGGGAAGGCTTGGAGGTGGTGTCTTGTGGCCCTCTATGGTGGTTAGGTAATTTCTC 1800  
 Qy 1801 AGAAGACAGACTGTGAAAATTTAGATAATGTCTGATGTCTATCATTTTCAATATACCAAAAA 1860  
 Db 1801 AGAAGACAGACTGTGAAAATTTAGATAATGTCTGATGTCTATCATTTTCAATATACCAAAAA 1860  
 Qy 1861 ACCCTGTGTCCGATGGCTATAAAGCAGCAACTCTGCTGCTCTCCCATCAAGAGCAGAG 1920  
 Db 1861 ACCCTGTGTCCGATGGCTATAAAGCAGCAACTCTGCTGCTCTCCCATCAAGAGCAGAG 1920  
 Qy 1921 ACACCTAAACAGGTAAGCACTCTACAGACCTCTACAGACAAATCATCTGTGTGGTACCATGCTA 1980  
 Db 1921 ACACCTAAACAGGTAAGCACTCTACAGACCTCTACAGACAAATCATCTGTGTGGTACCATGCTA 1980  
 Qy 1981 CCGAGCAAAATGCTCCCTGATGTTTGGCTTTTGGCTCTCTCCTCACTAAAGGCTCTCCT 2040  
 Db 1981 CCGAGCAAAATGCTCCCTGATGTTTGGCTTTTGGCTCTCTCCTCACTAAAGGCTCTCCT 2040  
 Qy 2041 CTCACTTATCAACTGTGTGACACTTGTGCGATCTCTGATGGCTGTCTGCGAATACTATG 2100  
 Db 2041 CTCACTTATCAACTGTGTGACACTTGTGCGATCTCTGATGGCTGTCTGCGAATACTATG 2100  
 Qy 2101 AGTTTTCCTTATGGGGAATTTGGCCGCGAGCTGCTGCTTCTCATTTGCCCTGTGGGCC 2160  
 Db 2101 AGTTTTCCTTATGGGGAATTTGGCCGCGAGCTGCTGCTTCTCATTTGCCCTGTGGGCC 2160  
 Qy 2161 CAGGAGCAAAATGGCTGCGCTCAACACCGGCTGCAAGCTTGAAGGTGTCCAACTTCCAG 2220  
 Db 2161 CAGGAGCAAAATGGCTGCGCTCAACACCGGCTGCAAGCTTGAAGGTGTCCAACTTCCAG 2220  
 Qy 2221 CAGCGGTACATCGTCAACCGCACCTTTATGTCGCCAAGGAGGTACAGCTGCTCTCTTT 2280  
 Db 2221 CAGCGGTACATCGTCAACCGCACCTTTATGTCGCCAAGGAGGTACAGCTGCTCTCTTT 2280  
 Qy 2281 CTCTCCATACCGCTTGGCAATTTTCTCTGAAGCACTTTCGAACTTTTGGGGGCGCTTTA 2340  
 Db 2281 CTCTCCATACCGCTTGGCAATTTTCTCTGAAGCACTTTCGAACTTTTGGGGGCGCTTTA 2340  
 Qy 2341 TCTCCGAGGCTCTCACTACCTATGTTTCTGTTCTCTTTAGAGACTCTTTTAAAGGACTGGGT 2400  
 Db 2341 TCTCCGAGGCTCTCACTACCTATGTTTCTGTTCTCTTTAGAGACTCTTTTAAAGGACTGGGT 2400

QY	2401	CTTTTCTATTCTATTCAAGGCTCAGAGCAATTTCTCTATCTTGGCCTTCAGGACACA	2460
Db	2401		
QY	2461	CTTTTCTATTCTATTCAAGGCTCAGAGCAATTTCTCTATCTTGGCCTTCAGGACACA	2460
Db	2461		
QY	2461	TATACTGAATTTTATCTACAGAGCGCATTTAGAAAGCCACCCAGACTGCAATACCTTTC	2520
Db	2461		
QY	2521	TATACTGAATTTTATCTACAGAGCGCATTTAGAAAGCCACCCAGACTGCAATACCTTTC	2520
Db	2521		
QY	2521	CATTTCTGTGTCTCTCTCTGAACCTCATCTCTTGTGGCTACTCTCTGAGACCCACTGCG	2580
Db	2521		
QY	2581	GACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTTGTCAACCCAGGCACTTAGGGT	2640
Db	2581		
QY	2641	TTTCTCTCTTTCAGGCGAGCTTGCAGATAAACAAACACAGACGTCCTGCTCATCGGGGAGA	2700
Db	2641		
QY	2701	AACCTGTTCCGAGGAGTCAGTGAAGTCTCTCACTGTGATGAGCAGGGCTAGCTCGGGGAGC	2760
Db	2701		
QY	2761	TGCTGACCCCTCTGGGATAGTCTGAGTATGACCCCTCTGTCTTCTTGTCTACTCTGAGG	2820
Db	2761		
QY	2821	CTAAAGATCAGTCTACCTGATCAAGCAGGTGCTCAACTTCCACCTGGAGAGCTTCTGC	2880
Db	2821		
QY	2881	TCCTCTTCTATTCCAGTAAGAACCGAGGTCCTGCCCTCTCTCTCTTCAAGAGTACGGA	2940
Db	2881		
QY	2941	TCAGCAATCAGCTCAGTCTCTGTGTGAAGTCTGACTCTGGCTACTATGCTCTCTCTTT	3000
Db	2941		
QY	3001	CTCTCTTCTATTCCAGTAAGAACCGAGGTCCTGCCCTCTCTCTCTTCAAGAGTACGGA	3060
Db	3001		
QY	3061	GGGCTTCAGCACCAACCATCATAGGCCACTTGAATAAGTCAAAAGCTTTTGGCTTC	3120
Db	3061		
QY	3121	AATTGAGTAATACCTTTGAGTTGTATGAGTGAAGCTTTATTTGTTTATCCATGGAAAGA	3180
Db	3121		
QY	3181	AATCAACTCAAAATCTGTAGGATGAGAAAGATGTTGGGAACGAAAGAGCCCTAGATAGA	3240
Db	3181		
QY	3241	GAAAACAGATCTGCTGAGTATAGTACTTATGGGGGAGCAGGGGGCGATATCCACTGAGTA	3300
Db	3241		
QY	3301	CAAGTACTTGTGGGAGAGAAATCCACTGAGTACAAAGTACTTGTGGCATGGAGATCCAC	3360
Db	3301		
QY	3361	TGAGTACAACTCTTGTGGGGAGGAGGATGGACAGAGCAAAAGTTGAAGGGAAGAG	3420
Db	3361		
QY	3421	ATGGAGAGGCTCATCTGGTTGGGGGTGTGAAAGGTCACTCTCTTTTCCATGTGATGGAGGT	3480
Db	3421		
QY	3481	TAAGAAAAACCAAGTGTGTGATTTGATGTCTTCAGACACCCCAACTATGAAACATATCC	3540
Db	3481		

Db	3481		TAAGAAAAACCAAGTGTGTGATTTGATGTCTTCAGACACCCCAACTATGAAACATATCC	3540
QY	3541		ACGAGAGCGGGCAGACTGTGGGAGACCTGGGCATTTAGGGAAGCGCGGCTTTTTCACAG	3600
Db	3541		ACGAGAGCGGGCAGACTGTGGGAGACCTGGGCATTTAGGGAAGCGCGGCTTTTTCACAG	3600
QY	3601		AGAAACTTTATGCTCATCTCTTGTGTCTACATCTCCCACTTTGTATGAGGTTTCACTCAGGT	3660
Db	3601		AGAAACTTTATGCTCATCTCTTGTGTCTACATCTCCCACTTTGTATGAGGTTTCACTCAGGT	3660
QY	3661		TTGCTTTCTACCGTTCTTCTGCTACTGCTGGAACCTTCAGTAGGATTTCCCAAGACGAGGA	3720
Db	3661		TTGCTTTCTACCGTTCTTCTGCTACTGCTGGAACCTTCAGTAGGATTTCCCAAGACGAGGA	3720
QY	3721		CAGCTCTTCTGTAAAGGAGGAGCCTCGATTTTCACTGTCTTAGAGAACTTCAGTAGGATTTCCCAAGACGAGGA	3780
Db	3721		CAGCTCTTCTGTAAAGGAGGAGCCTCGATTTTCACTGTCTTAGAGAACTTCAGTAGGATTTCCCAAGACGAGGA	3780
QY	3781		GAATCTAGGTCAAACGTAATCTTAGGTCAAGCGGGCAAAAATGACTGAACCGCTCTATT	3840
Db	3781		GAATCTAGGTCAAACGTAATCTTAGGTCAAGCGGGCAAAAATGACTGAACCGCTCTATT	3840
QY	3841		CCAGGTGAACGCTCAGCTGCTCAGATATCTAGAGTATTTGGGCTCCCAAGGATTAAGAT	3900
Db	3841		CCAGGTGAACGCTCAGCTGCTCAGATATCTAGAGTATTTGGGCTCCCAAGGATTAAGAT	3900
QY	3901		TCCTGTTAGTGAAGTCTCTTTTATTTTTCAGCAGCATCAGCGGTGACGACGAGCAATCCAG	3960
Db	3901		TCCTGTTAGTGAAGTCTCTTTTATTTTTCAGCAGCATCAGCGGTGACGACGAGCAATCCAG	3960
QY	3961		AAGAATGTCCAGAAAGGCTGAAGGAGACAGTGAAGGAGTACTATTGGCAAGCCCAATACT	4020
Db	3961		AAGAATGTCCAGAAAGGCTGAAGGAGACAGTGAAGGAGTACTATTGGCAAGCCCAATACT	4020
QY	4021		AAGCCATTCAGTAGAGAGCTGGGATTTCTTCTCTGCTTCCAGTCCCTTCTACTTTG	4080
Db	4021		AAGCCATTCAGTAGAGAGCTGGGATTTCTTCTCTGCTTCCAGTCCCTTCTACTTTG	4080
QY	4081		TAACATTTTATTTGACTGCTCTACTATCTGCTGCTCAATTTCTCGCTTAGCTGACCTGATC	4140
Db	4081		TAACATTTTATTTGACTGCTCTACTATCTGCTGCTCAATTTCTCGCTTAGCTGACCTGATC	4140
QY	4141		TAGCTGGGCTATAGATCTTTCAATCTGTCTTAAATTTTGAAGTCACAATTTCTGGAGCT	4200
Db	4141		TAGCTGGGCTATAGATCTTTCAATCTGTCTTAAATTTTGAAGTCACAATTTCTGGAGCT	4200
QY	4201		ACGAGAAAGCTTAGCTCAGCCAGTCTCATGAGCACTTGTCTCGAGGATGGCTTGTGACAG	4260
Db	4201		ACGAGAAAGCTTAGCTCAGCCAGTCTCATGAGCACTTGTCTCGAGGATGGCTTGTGACAG	4260
QY	4261		AGTCAATGTAGAAGACAGCATCCCTGATTTCCAGCTCTGCACTTCCCTAGTGGCCATGT	4320
Db	4261		AGTCAATGTAGAAGACAGCATCCCTGATTTCCAGCTCTGCACTTCCCTAGTGGCCATGT	4320
QY	4321		GTAATTTACTTTGGCTTTGATTAAGTATTTGGGAAAGCCAGTTCACGAGCTTACATAATC	4380
Db	4321		GTAATTTACTTTGGCTTTGATTAAGTATTTGGGAAAGCCAGTTCACGAGCTTACATAATC	4380
QY	4381		TGAAGAACCATGCAATTTGAAACTTAGAAAGCTGGGCAAAAATTTCTAGAGATGATTTTGG	4440
Db	4381		TGAAGAACCATGCAATTTGAAACTTAGAAAGCTGGGCAAAAATTTCTAGAGATGATTTTGG	4440
QY	4441		AGCTCAATTAACCGGATGCTCTGAAATGTGGCAAAAATCAACCCAGGATTAACAACAAAGAG	4500
Db	4441		AGCTCAATTAACCGGATGCTCTGAAATGTGGCAAAAATCAACCCAGGATTAACAACAAAGAG	4500
QY	4501		CTGGATTTTGCATAATAGGACAGTATTTAGAACTCAGTGTATTAATAGCTATCATCTTAAT	4560
Db	4501		CTGGATTTTGCATAATAGGACAGTATTTAGAACTCAGTGTATTAATAGCTATCATCTTAAT	4560
QY	4561		TAAATATATAGGCGCTATATATATATTTAAGATTAACAACAAGTGGATAGCTCCCAAT	4620
Db	4561		TAAATATATAGGCGCTATATATATATTTAAGATTAACAACAAGTGGATAGCTCCCAAT	4620

D	b	4561	TAAATATATAGGGCCTATATATATATATTATTAAGATTAAACAAGAGTGGATAGCCTCCCAAT	4620
Q	y	4621	TTACTTGCGCCTGGTTTTCAAAGAGTAGTAAAAATATACGTGATGGATTAAATTATATAGTCATG	4680
D	b	4621	TTACTTGCGCCTGGTTTTCAAAGAGTAGTAAAAATATACGTGATGGATTAAATTATATAGTCATG	4680
Q	y	4681	AAAGTATGAGATGGAAAACCCCTTTCCTTCTAATTTCTTTTACCTTCTTTAGTTTTTTTTTTC	4740
D	b	4681	AAAGTATGAGATGGAAAACCCCTTTCCTTCTAATTTCTTTTACCTTCTTTAGTTTTTTTTTTC	4740
Q	y	4741	TTCACACCCTGATCAAGGCCACTAGTAGACACCTATCTGCTGTGAGCTATTATATGACTTT	4800
D	b	4741	TTCACACCCTGATCAAGGCCACTAGTAGACACCTATCTGCTGTGAGCTATTATATGACTTT	4800
Q	y	4801	ACAGCAAAACAATTTGCTGTGTGCCTCTCTTTGGGAAGGAAACAGATAGCAGAGGCTC	4860
D	b	4801	ACAGCAAAACAATTTGCTGTGTGCCTCTCTTTGGGAAGGAAACAGATAGCAGAGGCTC	4860
Q	y	4861	AGGCTAGCAAGTCTGACTTTCCTTAAAGCCAGAGGCAATGTTGATAGCAGAGAAGTGAG	4920
D	b	4861	AGGCTAGCAAGTCTGACTTTCCTTAAAGCCAGAGGCAATGTTGATAGCAGAGAAGTGAG	4920
Q	y	4921	GCTCTTCGCAAGTGGGTGTGCTTAAGTAAGTACAGAAACAGAAAGGCTCCGGTGTAGTGAAT	4980
D	b	4921	GCTCTTCGCAAGTGGGTGTGCTTAAGTAAGTACAGAAACAGAAAGGCTCCGGTGTAGTGAAT	4980
Q	y	4981	TATCAGTAAGATATCTACCCCTTATCTCCTTCTATCGAACCTAAATCGTCTCTTTTCTTG	5040
D	b	4981	TATCAGTAAGATATCTACCCCTTATCTCCTTCTATCGAACCTAAATCGTCTCTTTTCTTG	5040
Q	y	5041	TGTTGAGGCTCATATAACACACTTCTTTTCTTTTGTAGTGTTTCATGGCTTTGTAGATTTTA	5100
D	b	5041	TGTTGAGGCTCATATAACACACTTCTTTTCTTTTGTAGTGTTTCATGGCTTTGTAGATTTTA	5100
Q	y	5101	GTGCTCTGCCAGTCTTGTGTAGAGGTTTGTACTTTCAGACACCTGGGCTTGGATGTAGC	5160
D	b	5101	GTGCTCTGCCAGTCTTGTGTAGAGGTTTGTACTTTCAGACACCTGGGCTTGGATGTAGC	5160
Q	y	5161	ATGCCAAGGCACACACTTCCTGTAATGCTGTGTAAAGGTTATTATTCACTTTACTTTGTC	5220
D	b	5161	ATGCCAAGGCACACACTTCCTGTAATGCTGTGTAAAGGTTATTATTCACTTTACTTTGTC	5220
Q	y	5221	TTTGGAAAGGTGAACGCTGTGTGAGAAGAACTCACAGGAGATGTGTCTCTGTAGGAAA	5280
D	b	5221	TTTGGAAAGGTGAACGCTGTGTGAGAAGAACTCACAGGAGATGTGTCTCTGTAGGAAA	5280
Q	y	5281	ACTTTTTTTTTCCCCTTAAATGCTTAATTCACCTTCCTCAGTCAACTTTGACTTTTATACC	5340
D	b	5281	ACTTTTTTTTTCCCCTTAAATGCTTAATTCACCTTCCTCAGTCAACTTTGACTTTTATACC	5340
Q	y	5341	ATGCTGTACATGAAGAGTGTATTAGGCCGCTCTCATGGCTCTGGGAAAAGCACCAATA	5400
D	b	5341	ATGCTGTACATGAAGAGTGTATTAGGCCGCTCTCATGGCTCTGGGAAAAGCACCAATA	5400
Q	y	5401	GGGGAAGGAATGTTATGCTGAGAAATCTGACCGGCAGGGAACCTGGTCAGAGCTCCCCCG	5460
D	b	5401	GGGGAAGGAATGTTATGCTGAGAAATCTGACCGGCAGGGAACCTGGTCAGAGCTCCCCCG	5460
Q	y	5461	AAGACCAACACAGGTGTTAAGTAGGAAACAGTCCAGGTGGGCTCATGTAAATAGAAATGAA	5520
D	b	5461	AAGACCAACACAGGTGTTAAGTAGGAAACAGTCCAGGTGGGCTCATGTAAATAGAAATGAA	5520
Q	y	5521	CAGAGGGAGGAAGATAGCTACAAAGTTTTCATAGGTCCTGGAGCTTTAAAGATACAAA	5580
D	b	5521	CAGAGGGAGGAAGATAGCTACAAAGTTTTCATAGGTCCTGGAGCTTTAAAGATACAAA	5580
Q	y	5581	TAGCTGCTTGGGCTTCATAAACAAAGGAAGTCTGGGAAGGCAGCAAGTGAGAGGAAATGG	5640
D	b	5581	TAGCTGCTTGGGCTTCATAAACAAAGGAAGTCTGGGAAGGCAGCAAGTGAGAGGAAATGG	5640
Q	y	5641	AAAGGGAAAAACAGAAATGTAGAGGACTTGAACAGCTACAAATCCTCTTACAGACGATTT	5700
D	b	5641	AAAGGGAAAAACAGAAATGTAGAGGACTTGAACAGCTACAAATCCTCTTACAGACGATTT	5700

Qy	5701	TTCTTTGGAA	CAATCTAGAA	GGTGTG	TGATTTAG	CTGATTTG	CAAGGGG	GAC	TGCTTTTG	CGCAT	5760							
Db	5701	TTCTTTGGAA	CAATCTAGAA	GGTGTG	TGATTTAG	CTGATTTG	CAAGGGG	GAC	TGCTTTTG	CGCAT	5760							
Qy	5761	TTGAA	CTCTGG	TTTTTGT	CTCTCC	ATTTAG	AGTTTGA	AAAGCG	CTCAC	CCCTTTT	TATAC	CCCTCGAA	5820					
Db	5761	TTGAA	CTCTGG	TTTTTGT	CTCTCC	ATTTAG	AGTTTGA	AAAGCG	CTCAC	CCCTTTT	TATAC	CCCTCGAA	5820					
Qy	5821	TGGAG	GAGG	AAAG	AGGG	GTGT	TATAG	CTCT	CACTCG	TGG	AGTTT	TAC	TAG	TTTAC	GCA	ATG	5880	
Db	5821	TGGAG	GAGG	AAAG	AGGG	GTGT	TATAG	CTCT	CACTCG	TGG	AGTTT	TAC	TAG	TTTAC	GCA	ATG	5880	
Qy	5881	GAAC	GAG	CAC	ACTCGG	ACCTCCT	TTG	ACAAAA	AAAAAT	TG	GAAC	CTGTG	TTTCT	GTCT	TTG	TTT	5940	
Db	5881	GAAC	GAG	CAC	ACTCGG	ACCTCCT	TTG	ACAAAA	AAAAAT	TG	GAAC	CTGTG	TTTCT	GTCT	TTG	TTT	5940	
Qy	5941	TGTT	CTTTTT	TGTT	TAAG	AAAG	CAC	AGG	CAAA	CCCG	ACC	AC	CAC	ATG	GGT	TGA	AT	6000
Db	5941	TGTT	CTTTTT	TGTT	TAAG	AAAG	CAC	AGG	CAAA	CCCG	ACC	AC	CAC	ATG	GGT	TGA	AT	6000
Qy	6001	GAGT	CA	AGG	CTTT	TG	AGT	GAG	CA	CTCAT	CA	ATAG	TTG	AT	CAT	GGT	CA	6060
Db	6001	GAGT	CA	AGG	CTTT	TG	AGT	GAG	CA	CTCAT	CA	ATAG	TTG	AT	CAT	GGT	CA	6060
Qy	6061	CCTGT	C	AGG	CCG	AG	CCCTG	CTG	CG	CTT	CG	CA	CTT	CA	AGT	CT	C	6120
Db	6061	CCTGT	C	AGG	CCG	AG	CCCTG	CTG	CG	CTT	CG	CA	CTT	CA	AGT	CT	C	6120
Qy	6121	CTG	CT	CTT	TAG	CAC	AGT	TAG	GAGT	TG	GAG	CAAA	CCCTTTTT	T	T	T	T	6180
Db	6121	CTG	CT	CTT	TAG	CAC	AGT	TAG	GAGT	TG	GAG	CAAA	CCCTTTTT	T	T	T	T	6180
Qy	6181	AA	TTG	CA	AAA	AG	CTGT	GA	TAA	TTTGG	GAT	TA	AT	TG	AT	CT	AT	6240
Db	6181	AA	TTG	CA	AAA	AG	CTGT	GA	TAA	TTTGG	GAT	TA	AT	TG	AT	CT	AT	6240
Qy	6241	TGT	CG	AA	GGT	CA	AT	AG	AT	AG	AT	TA	AT	AG	CC	CA	T	6300
Db	6241	TGT	CG	AA	GGT	CA	AT	AG	AT	AG	AT	TA	AT	AG	CC	CA	T	6300
Qy	6301	CA	AGT	TA	AT	AG	TAG	TGC	CTGT	GGT	GT	CC	TT	AG	GT	C	A	6360
Db	6301	CA	AGT	TA	AT	AG	TAG	TGC	CTGT	GGT	GT	CC	TT	AG	GT	C	A	6360
Qy	6361	TTGG	CA	AA	T	AT	TAC	TG	CT	GT	CA	AA	AT	TAC	TG	CT	GT	6420
Db	6361	TTGG	CA	AA	T	AT	TAC	TG	CT	GT	CA	AA	AT	TAC	TG	CT	GT	6420
Qy	6421	AG	AG	GC	T	GA	CT	TGG	TTTT	TGG	TG	CT	CAG	CA	AG	CA	AT	6480
Db	6421	AG	AG	GC	T	GA	CT	TGG	TTTT	TGG	TG	CT	CAG	CA	AG	CA	AT	6480
Qy	6481	GT	AC	CA	CT	TT	AG	AA	T	CT	AC	CT	GT	CA	AA	T	CT	6540
Db	6481	GT	AC	CA	CT	TT	AG	AA	T	CT	AC	CT	GT	CA	AA	T	CT	6540
Qy	6541	TGG	G	AG	AG	TG	G	AG	AT	CA	AG	CG	AT	T	GGG	GA	CT	6600
Db	6541	TGG	G	AG	AG	TG	G	AG	AT	CA	AG	CG	AT	T	GGG	GA	CT	6600
Qy	6601	AA	AT	CT	CT	TG	CG	T	CT	CAG	CG	AG	AA	GA	AG	CT	T	6660
Db	6601	AA	AT	CT	CT	TG	CG	T	CT	CAG	CG	AG	AA	GA	AG	CT	T	6660
Qy	6661	TAAA	AG	AA	CA	AT	AG	AT	CC	CT	GT							



[illegible]

2276	Db	TTGGCAAGCCACAATACTAAGCCATTCAGTAGAGAGCGTGGGATTTCTTTCTCTCGCTTC	2333
4063	Qy	CCAGTCCCTTCTACTTTTGTAAACATTTTATTGTGCTTGTCTACTATCTGCGPCCAACTACTCG	4122
2336	Db	CCAGTCTCTTCTACTTTTGTAAACATTTTCTTTGACTTGTCTACTGTCTGGTCCCACTACTCA	2395
4123	Qy	CTTAGCTGCACTGTATCTAGCTGGGTCTATAGATCTTTTCAATCTGTGTCTTAAATTTGTGA	4182
2396	Db	CTTAGCTGCACTGTGCATCTAGCTGGGTCTATAGATCTTTTCAATCTGTGTCTTAAATTTGTGA	2455
4183	Qy	AGTCACAATTTCTGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTCATATGACACTTGTCTCG	4242
2456	Db	AGTCACAATTTCTGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTCATATGACACTTGTCTCG	2515
4243	Qy	GAGGATGGCTTGTGACAGAGTCAATGCTAGAAGACAGCATCCCTGATTTCCAGCTCTGCA	4302
2516	Db	GAGGATGGCTTGTGACAGAGTCAATGCTAGAAGACAGCATCCCTGATTTCCAGCTCTGCA	2575
4303	Qy	CTTGCTTAGTGGCCATGTGTAAATTACTTTTGGCTTGTATTAAGTATTTTGGGAAAGCCAGTTC	4362
2576	Db	CTTGCTTAGTGGCCATGTGTAAATTACTTTTGGCTTGTATTAAGTATTTTGGGAAAGCCAAATTC	2635
4363	Qy	CCACGGACTACATAATCTGAAGAACCATGCAATTGAAAACATGAAAAGCTGGGCACAAACT	4422
2636	Db	CCACGGACTACATAATCCGAAGAAGCATGCAATTGAAAACATGAAAAGCTGGGCACAAACT	2695
4423	Qy	TACTAGAGATGATTTTTTGGACTCAATTAACCGGATGCTCTCAAAATGTGGCAAAATCAACCC	4482
2696	Db	TACTAGAGATGATTTTTTGGACTCAATTAACCTGATGCTCTGAAATGTGATCAAACTCAACCC	2755
4483	Qy	AGAATAACAACAAAGAGCTGGATTTTGCAAAATAGGACAAGTATTTAGAATCATCTGGTATTT	4542
2756	Db	AGAATAACAACAAAGAGCTGGATTTTGCAAAATAGGACAAGTATTTAGAATCATCTGGTATTT	2815
4543	Qy	AATAGCTTATCATCTTAATTAATAATATAGGSCCTATATA----TATATTTAAGATTAACA	4598
2816	Db	AACAGCTGTATCTTAAATTAATAATATAGTGTCTATTTAGCTGCTATTTAAGATTAACA	2875
4599	Qy	CAAGAGTGGATAGGCTCCCAATTTACTTTGGCTGGTTTCAAAAGAGTAAAAATATCAGTC	4658
2876	Db	CAAGAGTGGATTAACCTTCCCAATTTACTTGGGCTGGTTTCAATAGAGTAAAAATATCAGTC	2935
4659	Qy	ATGGATTAATTTATAGTGTCAATGAAGATGAGATGGAAACCTTTTCCCTTACTTTTTTAACTT	4718
2936	Db	ATAGATTAATTTATAGTGTCAATGAAGATGAGATTTGGAAACCTTTTCCCTTACTTTTTTAACTT	2995
4719	Qy	TCATTTCTTAGT-----TTTTTTTTTCTTACACCCCTGATCAAGCCACTAGTAGCACCTT	4773
2996	Db	TCATTTCTTAGT-----TTTTTTTTTCTTACACCCCTGATCAAGCCACTAGTAGCACCTT	3055
4774	Qy	ATCTGCTGTGAGCTATTATATGACTTTTACAGCAAAACAATGCTGTGTGGCCCTCTTTGG	4833
3056	Db	ATCTGCTGTGAGCTATTATATGACTTTTACAGCAAAACAATGCTGTGTGGCCCTCTTTGG	3115
4834	Qy	GGAAAGGAAACAGGATAGCAGGAGGCTCAGGCTAGCAAGTCT-GACTTGGCCCTAAAGCCAG	4892
3116	Db	GGAAAGGAAACAGGATAGCAGGAGGCTCAGGCTAGCAAGTCTGAGACTCAACCTAAAGCCAG	3175
4893	Qy	AGGCAATGGTTGATAGCAGAAAGTGAAGGCTCTTTCGCAAGTGGGTGTGCTTTAAGTAATCA	4952
3176	Db	AGGCAATGGTTGATAGCAGAAAGTGAAGGCTCTTTCACAAAGTGGGTGTGCTTTAAGTAATCA	3235
4953	Qy	GAACACGGAAGGCTCCGGTTGATGGAATTAATCAGTAAGATATCTACCCCTATCTC---CT	5009
3236	Db	GAACACGGAAGGCTCTGGTTGATGGAATTAATCAGTAAGATATCTACCCCTATCTCCTTCT	3295
5010	Qy	TCTATCGAACCTAAATCGTCTCTTTTCTTGTGTGTAGGCTGATAAACAACACTTTGTTTC	5069
3296	Db	TCTATAGAAAGCTAAACCGTCTCTCTTCTTGTGTGTAGGCTGATAAACAACACTTTGTTTC	3355
5070	Qy	TTTTGAGTGTTTCATGGCTTTGTAGATTTTATAGTGTCTCTGCCAGTCTTGTGTAGAGGTTT	5129
3356	Db	TTTTGAGTGTTTCATGGCTTTGTAGATTTTATAGTGTCTCTGCCAGTCTTGTGTAGAGGTTT	3415



QY 5130 GTTACCTTGCACCTGGCTTGGATGTTAGCATGCCAAAGGCACACACTTCTGAATGCCT 5189  
DB |||||  
3416 GTTACCTTGCACCTGGCTTGGATGTTAGCATGCCAAAGGCACACACTTCTGAATGCCT 3475  
QY |||||  
5190 GTGTAAAGGTTATTTATCAATTTTACATTTTGTCTTTGGAAAGGTGAAGCGTGTGTGAGAAAG 5249  
DB |||||  
3476 GTGTAAAGGTTATTTATCAATTTTACATTTTGTCTTTGGAAAGGTGAAGCGTGTGTGAGAAAG 3535  
QY |||||  
5250 AACTCACAGGAGATGTCTCTGTAGTAAACCTTTTTCCTTAAATGCCTATAA 5309  
DB |||||  
3536 AACTCACAGGAGATGTCTCTGTAGTAAACCTTTTTCCTTAAATGCCTATAA 3594  
QY |||||  
5310 TCACATTTTACAGTCAACTTTTACATTTTATACCATGTCTCACATGAAGAGTGTTTAGGCC 5369  
DB |||||  
3595 TCACATTTTACAGTCAACTTTTACATTTTATACCATGTCTCACATGAAGAGTGTTTAGGCC 3654  
QY |||||  
5370 CGCTCTCATGCTCTGGGAAAGCAACATAGGGGAAGGATGTTATGCTGAGAAATCTG 5429  
DB |||||  
3655 CGCTCTCATGCTCTGGGAAAGCAACATAGGGGAAGGATGTTATGCTGAGAAATCTG 3714  
QY |||||  
5430 ACCGGCAGGAAACTGTCTAGAGCTCCCGAAGACCAACAGGCTGTTAAGTAGGAACA 5489  
DB |||||  
3715 ACTGGCAGGAAACTGGCTCAGAGCTCCCAAGACCACTACAGGTGTTAAGTAGGAACA 3774  
QY |||||  
5490 GTCCAGGGTGGCTCATGTTAATGAATGGAACAGAGCGAGGGAAGATTAAGCTACAAAGTT 5549  
DB |||||  
3775 GTCCAGGGTGGCTCATGTTAATGAATGGAACAGAGCGAGGGAAGATTAAGCTACAAAGTT 3834  
QY |||||  
5550 TCATAGGGTCCGAGTCTTAAAGTACAAATAGCTGCTTGGCTTCATTAACAAAGGAAG 5609  
DB |||||  
3835 TCATAGGGTCCGAGTCTTAAAGTACAAATAGCTGCTTGGCTTCATTAACAAAGGAAG 3894  
QY |||||  
5610 TCTGGGAAGCAGCAAG--TGAGAGGGAATGGAAGGGAAGGAAACAGAGATGTAGAGGA 5666  
DB |||||  
3895 TCTGGGAAGCAGCAAGTGTAGAGGGAAGTGGGAAGGGAAGGAAAC--AATGTAGAGGA 3952  
QY |||||  
5667 CTTGGAACAGCTACAAATCTCTACAGACGATTTTCTTGGAACAATCTAGAA--GGT 5722  
DB |||||  
3953 TTTGGAACAGCTACAAATCTCTACAGAGGATTTTCTTGGAACAATCTAGAACAGGGT 4012  
QY |||||  
5723 AGTGGAATAGGT--GATGCGAGGGGACTGTGTTTGCATTTGAATCTGGGTTTTTGTCTC 5781  
DB |||||  
4013 GGTGGATTTAGGTGATCGCAGAGGACTGTGTTTGCATTTGAATCTGGGTTTTTGTCTC 4072  
QY |||||  
5782 TCCATTGAGGTGGAAGCGTCAACC--TTTTTACCTCGAATGGAGGGAAGAGCGGT 5840  
DB |||||  
4073 TCCATTGAGGTGGAAGCGTCAACCCTTTTACCTCGAATGGAGGGAAGAGCGGT 4132  
QY |||||  
5841 GTTATGACTCTTACCTGGAGTTTACTAGTTTACGCAATGGAACAGACACTCGGGACCTC 5900  
DB |||||  
4133 GTTTTGNCTCTACCTGGAGTTTACTAGTTTACGCAATGGAACAGACACTCGGGACCTC 4192  
QY |||||  
5901 CTCTTGAC-----AAAAAATGGAACCTGTGTTTGTCTGTTTGTCTTTGTTGTTG 5950  
DB |||||  
4193 CTCTTGACAGAAAAAAGGAAAGGAAAGGAAAGGAAAGGAAAGGAAAGGAAAGGAAAGG 4252  
QY |||||  
5951 TTAAGAAAGCACA----- 5963  
DB |||||  
4253 TTAAGAAAGCAGGCGAGCTGGGCAATGCTGGGCAATGCTTTTAAATCCAGCATTTGGGAG 4312  
QY |||||  
5964 ----- 5963  
DB |||||  
4313 GCAGGCGAGGTGACTTTCTAAATTTCAAGCGCAGCTGGTCTACAAAGTGAAGTTCCAGGA 4372  
QY |||||  
5964 ----- 5963  
DB |||||  
4373 CAGCCAGGGCTATACAGAGAAACCTGTCTCGGGAAGGAAAGGAAAGGAAAGGAAAGG 4432  
QY |||||  
5964 ----- 5963  
DB |||||  
4433 AAAAGAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 4492

QY 5964 ----- 5963  
DB |||||  
4493 GAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 4552  
QY |||||  
5964 ----- 5963  
DB |||||  
4553 GAGAAGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 4612  
QY |||||  
5964 ----- 5963  
DB |||||  
4613 AAGAGAAAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 4672  
QY |||||  
5986 TGAATGTGGTCTTTGAGTCAAGGCTTTTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGA 6045  
DB |||||  
4673 CGTATGTGGTCTTTGAGCAAGGCTTTTGAATGAGCGCTCATCAATAGTGTGATCATGG 4732  
QY |||||  
6046 TCAGGTGAGGGGTACTGTCTGAGCGGCGGCTTGGCTTCGCACTTAACATCTCCAGG 6105  
DB |||||  
4733 TCAGGTGAGGGGTACTGTCTGAGCGGCGGCTTGGCTTCGCACTTAACATCTCCAGG 4792  
QY |||||  
6106 TCTCAGTATCACTTCTCTGCTTACTTAGCACAAGTGTAGGAGTTGAGCAAACTTTTTCCTCA 6165  
DB |||||  
4793 TCTCAGTATCACTTCTCTGCTTACTTAGCACAAGTGTAGGAGTTGAGCAAACTTTTTCCTCA 4852  
QY |||||  
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DB |||||  
4853 CCCCCAATAAATTGAACAAAAGGAGTGTGTAATTTTGGGATACAGTGTGATAATT 4912  
QY |||||  
6226 GATCTATGTGCTGTCGACGAGTTCATATAGATAGTATTAATAGGCCCACTCAACAGCTT 6285  
DB |||||  
4913 GATCTATGTGCTGTCGACGAGTTCATATAGATAGTATTAATAGGCCCACTCAACAGCTT 4972  
QY |||||  
6286 TATGGGTGTGAATGCAAGTAATATAGGTAGATGCTGCTGCTTCTAGGTGAGAAAGG 6345  
DB |||||  
4973 TATGGGTGTGAATGCAAGTAATATAGGTAGATGCTGCTGCTTCTAGGTGAGAAAGG 5031  
QY |||||  
6346 CATGATTTTAAAGTCTTGGGCAAAATCATATTAATCTCATGCTAAAAATACATTATGTGA 6405  
DB |||||  
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QY |||||  
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DB |||||  
5092 TTATCAATCTTTTAGAAGGCTGATCTTGGTTTTGGTCTCAGCAAGCAAAATGTCACC 5151  
QY |||||  
6466 AGCTCTTTCTAACTGTGTACCTTTAGAAAACTCTACCTGTGTCTCAAAATGGTTGTATT 6525  
DB |||||  
5152 AGCTCTTTCTAACTGTGTACCTTTAGAAAACTCTACCTGTGTCTCAAAATGGTTGTATT 5211  
QY |||||  
6526 CTTATTTTCTAACTGTGTACCTTTAGAAAACTCTACCTGTGTCTCAAAATGGTTGTATT 6585  
DB |||||  
5212 CTTATTTTCTAACTGTGTACCTTTAGAAAACTCTACCTGTGTCTCAAAATGGTTGTATT 5271  
QY |||||  
6586 GTTTATGTCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAAAGGAGAACTG 6645  
DB |||||  
5272 GTTTATGTCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAAAGGAGAACTG 5331  
QY |||||  
6646 CTCTTTCTGCTCTTAAAGAAACAAATGAATGCTTGAATGCTTTTTTACTTAAAGGA 6705  
DB |||||  
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QY |||||  
6706 AAGTGAGAGCTAAAGTCTCATCATATTAGAAGATTTCATGAAACCTGGCTCAGTTGA 6765  
DB |||||  
5392 AAGTGAGAGCTAAAGTCTCATCATATTAGAAGATTTCATGAAACCTGGCTCAGTTGA 5451  
QY |||||  
6766 AAAAGAAATAGTGTCAAGTTGCTCATGAGACGAGAGGTAGACTTGAATACCAAGAT 6825  
DB |||||  
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QY |||||  
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DB |||||  
5512 TCATTGACAAATATTTTATTTGTCTCATGATACACAGAAAAAATGACTTTTAAAAA 5571  
QY |||||  
6886 TTGTTTGAAGGAGGTTTACCTCTCATTTCTTTAGAAAAAAGCTTATGTAATCTTCTTC 6945

Db 5572 TTGTTTGAAGAGGTTACCTCTCATTTCTCTAGAGAAAGGCTATGTAACCTTC 5631  
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Db 5632 CATAAACCAATACTTTATATATAGTAAGTTATTTATTAATAAGTATACATTTTATATGTC 5691  
Qy 7006 AGTTTATATATAGGATTTATTTATAGAAACATTTATCTGCTATTTGATATTT-AGTATAAG 7064  
Db 5692 AGTTTATATATAGGATTTATTTATAGAAAAATTTATCTGATTTGATATTTGAGTATATA 5751  
Qy 7065 GCAATAATATTTATGACATAAATCTATGGAAACAAAGATATCTTAGGCTTTTAATAAACA 7124  
Db 5752 GCAATAATATTTATGATATAAATCTATAGAAACAAAGATATCTTAGGCTTTTAATAAACA 5811  
Qy 7125 TGGATATCATAAATCTTCTGTCTTGTATATTTTCTCCCTTTAATATCAACAATACCATCA 7184  
Db 5812 TGAATATCATAAATCTTCTGTCTTGTATATTTTCTCCCTTTAATATCAACAATACCATCA 5871  
Qy 7185 TCATCATCATTTACCAATCATCTCATGATTTTCATGCTTGACCCATATTTACTGTTAAA 7244  
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Qy 7245 GTTG 7248  
Db 5932 TTIG 5935  
RESULT 3  
US-11-177-987-26  
; Sequence 26, Application US/11/177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
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; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 26  
; LENGTH: 4797  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-11-177-987-26  
Query Match 9.2%; Score 686; DB 7; Length 4797;  
Best Local Similarity 53.8%; Pred. No. 3.7e-157;  
Matches 2644; Conservative 0; Mismatches 1875; Indels 393; Gaps 44;  
Qy 2034 CTCCTCTCACTTATCAACTGTGTGACACTGTGTGCGATCTCTGATGGCTGTCTGCAGAA 2093  
Db 29 CTCCTCTCCCAAGTACCAAGTTGCTGAGTTAGAAATTTGCTGCAATGGCCGCCCTGCAGAA 88  
Qy 2094 ATCTATGAGTTTTCCTTATGGGACATTTGGCCGCCAGCTGCCTGCTTCTCATTTGCCCT 2153  
Db 89 ATCTGTAGCTTTTCTTATGGGACCTTGGCCGCCAGCTGCCTTCTTCTTTGGCCCT 148  
Qy 2154 GTGGCCCGAGGAGGAATAGCGTCCCGTCAACACCCGGTGCAGCTGTAGGTGTCAA 2213  
Db 149 CTTGGTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGCGAGGCTTGCAGAGTCCAA 208  
Qy 2214 CTTCCAGCAGCGTACATCGTCAACCGCACCTTTTATGCTGGCCCAAGGAGGTACAGCTGCA 2273

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Qy 2274 TCTCTTTCTCTCATACCGCTTGGCATTTTCTCTGAGACACTTGCATACTTTTGGGG 2333  
Db 269 AATCTGCTCTTTCTCGTTGGATCTATTTGGAAATCCAAATAGTTCTTTAAACTTTTCTTCA 328  
Qy 2334 CGCTTTATCTCCGAGGTCTCACTACTATGTTT-----TCTGTCTCTTTAGAG 2382  
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Qy 2383 ACTCTTTAAGAGCTGGGTCTTTTCTATTTTCAAGGTCTCTCAGGACCATTTCTCTAT 2442  
Db 389 TTTTTCAGAGACTCTTTTGGGAATCTGGCTTTTTTTTTTCTTGAACCTCTTCTCCCTCAT 448  
Qy 2443 CTTGGGCTTCAGACACATATATGAATTTTATCTAAGAGGCTCTCAGGACCATTTTCC 2500  
Db 449 TTTGGGCTTTATGATACATATGATGAATTTTCCCAAGAGCGGCCCATTCAGTAATCCAT 508  
Qy 2501 CCCAGCACTGCAATCTTTCCATTTCTCTGCTCTCTCTGAACTCATACTCTCTTGGC 2560  
Db 509 CTGATGATTTTTTTTCTTTATGCTCTGCTGATTTTCTTAACTCATGACACATCTG 568  
Qy 2561 TACTC-----CTGAGACCCACTGCGGACATACATCTCTAC 2595  
Db 569 AATCTCTTTTGTATGATTTGCTCTGGGAGACGGGATGGGCACATGTCTAT 628  
Qy 2596 TTAAGGCTTTTCTCATCTCTGTCACCGAGGCACTTAGGGTTTC-TCTCTTTTCTAG 2654  
Db 629 GTATAAATTTTTTCTTTATTTGCTCAATGTCCAGACCTTTAGTCTTTTCTCTCCAG 688  
Qy 2655 GCCAGCCTTCAGATACACACAGACGTCGCGCTCATCGGGAGAACTTGTCCGAGGA 2714  
Db 689 GCTAGCTTGGCTGATACACACAGACGTCGCTCTCATTTGGGAGAAACTGTTCACGGA 748  
Qy 2715 GTCAGTCTAAGTCTCTCACTGTGATGACAGGCG-----TAGCTGCGGAGCT 2761  
Db 749 GTCAGTCTAAGTCTCACTGTGACGACGCGCGTGTGCGCTCATCGGTACTTTGGGT 808  
Qy 2762 GTTGGACCTCTGCGGATAG-----TCTGACGATGACCCCTGCTGCTTCTGTCTACCTGC 2817  
Db 809 GGTGCTGATGATGGTTTAGGTCTTATCCCTTATGACCCCTTTCTTTTCCCTTCCACCTGC 868  
Qy 2818 AGCTAAAGATCAGTCTACTGATGAAGACGAGTCTCAACTTCCACCTGGAAGAGTTC 2877  
Db 869 AGATGAGTGAGCGCTGCTATCTGATGAAGACGAGTCTGAACTTCCACCTTGAAGAGTGC 928  
Qy 2878 TGCTCCCTCCAGTCAGACAGGTTCCAGCCCTACATGACGAGGAGTGTACCTTTCTGACCA 2937  
Db 929 TGTTCCTCTCAATCTGATAGGTTCCAGCTTATATGACGAGGAGTGTGCTTCTGCGCA 988  
Qy 2938 AACTCAGCAATCAGCTCAGCTCCTGTGTAGTCTGACTCTGGGCTACCTATGCTCTCTCT 2997  
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Qy 2998 CTTCTCTTCTTATCCAGTAAGAACCCGAGTCTGCGCTCTCTCTCTTCAACAAGATGA 3057  
Db 1049 CTCCTTCTCTCTTCCACAGAGACCCCTTATACCCAACTCTCTCTCTCTTCCCTTACCCC 1108  
Qy 3058 GGAGGGCTCAGCACACCACCATCATAGGCCACTTGAANAATAGGTTCACAAAGGCTTTGGC 3117  
Db 1109 TAAGCTAGCAGGAAGAGTGTCTTGGCAGCAGTGTATCAGGAGTCA-----TTTGGG 1161  
Qy 3118 TTCAATTTGAGTAATACCTTTGAGTTTGTATGAGTGAAGCTTTATTTGTTTATCATGAA 3177  
Db 1162 ATCATAGAGTATTTGCTTTTGTGCTTGTAGTCACTCTTGTAGTTTATAGTGTGTAATG 1221  
Qy 3178 AGAATCACTCAATTTCTGTAGGATGAGAAAGATTTGGGAAACGAAAGAGGCTAGAT 3237  
Db 1222 GGGTCTGGAACTTAAGTGTACAGAGCGGCAATTTGGTTTGTCTTTCGAAAAAGGCACTC 1281  
Qy 3238 AGAGAAACAGATCTGCTGAGTATAGTATTATGCGGGGAGAGGGGGCGATATCCACTGA 3297



1282 A-----GGTTGCGTAAGATGAGAAAGGTTGG 1309 Db  
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1310 GAAAAATCTAGCTGTGGAAATGGATCCATTTAGTCTAAGTCTTGTGGGGGAGGGATGG 1369 Db  
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1370 CATGGAGAGAAATTTAGAAGAGAAAGTGGGAAATGGGAAGGCTTTAAA----- 1415 Db  
3418 AAGATGGAGAGGCTCATGTGTCGGGGTGTGAAGGTCACTCTTTTCCATGTGATGGAG 3477 Qy  
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3478 AGTTAAGAAAAACCA-GTGTGTGAGTTTGTATGTCTTCAGACACCCCACTATGAACAT 3536 Qy  
1461 AGCCACAAATCGAGGCGTGTGACTTGTATGCGCGTGAACATTTGAACTATGNAAAA 1520 Db  
3537 ATCCAGAGAGCGGAGACTGTGGGAGACCTGGGCAATTTAGGGAAGGCGC--GGCTTTT 3594 Qy  
1521 AGTTTGAGTGGAGTGGGCCCAAGAAAGGCCCTAGGACTTACTGAAGAGGCGCTTAATTTT 1580 Db  
3595 CACACAGAAACTTTATGCTCATCTCTGTGCTACACTCCCACTTTGATGAGGTTTCAGC 3654 Qy  
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3754 GTGTCCTAGAGAAAGTACTCAGAGAAATCTAGTCAAGTGAATCTAGTCAAGC 3813 Qy  
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3814 GGGCAAAATGACTGAAGCGCTCTATTCAGAGTGAACGGTCACTGCTCAGATATACTG 3873 Qy  
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3874 AGGTATTTGGCTCCCAAGGATGAATTTCTGTAGTGA-GTCTGCTTTTATTTTGCAGCA 3932 Qy  
1881 GAGTTTCATAAAGTTTCAGCACAAATTTAAGATAGTTATGCTGTTATTTATAGCA 1940 Db  
3933 CATTACGGGTGACGACAGAAATCTCAGAAATGTGCAAGGCTGAAGGAGACAGTGAA 3992 Qy  
1941 TATTGAAGGTGATGACCTGCATATCCAGAGGAATGTGCAAAAGCTGAAGGACACAGTGAA 2000 Db  
3993 AAGGTTACTATTGGCAAGCCACAAATCTAAGCCATTCAGTAG--GAGACGTGGGGATTC 4050 Qy  
2001 AAGGTTAGGACTGATACTGCTCAATGCTAAGTCAATGCAATAGGAGACAAATTTGTTT 2060 Db  
4051 TTTCTCTGCTCCAGTCCCTTCTACTTTGTATCAATTTTATTTGACTTTGTCTACTATCTG 4110 Qy  
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4111 GTCCATTTACTGCTTACTGCACTGTATCTAGCTGGGTCTATAGATCTTTTCAATCTGTG 4170 Qy  
2121 GGGCATTA-----CTTTGGTGTGTGTATGTAGATATATCTATATCTAGATGTAGTT 2176 Db  
4171 TCTAAATTT--GTAAGTCAATTTCTGGAGCTAGCAGAAAGCTTAGCTCAGCCGCTCTC 4227 Qy  
2177 TCCAAATCTTGGCAATTTGTAGAATTTCTAGAATCTGTTGGGATCTTAGCTTGTCTAGTCA 2236 Db  
4228 ATGAGCACTTGTCTGGAGGATGGCTGTGACAGAGTCAATGCTAGAGACAGCATCCCTG 4287 Qy  
2237 ATACCTCAGATTCGGGGATGGTCACTGGCAGAGATAGGGCTAGAAATGCAAGTCTCTG 2296 Db  
4288 ATTCCAGCTCTGCAC--TTGCCCTAGTGGCCATGTGTAATTTACTTTGGCTTTGATTAAGTAT 4346 Qy  
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4347 TTGGGAAA--GCCAGTTCCCGAGGACCTACATAATCTGAAGAACCAATGCAATGAAAACCTA 4404 Qy  
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4405 GAAAGCTGGGCA-----CAAACTTACTAGAGATGATTTTTGAGCTCAATTAACGAGTGCTC 4460 Qy  
2417 GAAAATGCTATGGGCAATTTATTTGAAGTCAATTTTTGAAGTCAATTAATGCAATGCTT 2476 Db  
4461 TGAATGTGGCAAAATCAACCCAGATAACAACAAGAGAGCTGGATTTGCAATAGSACA 4520 Qy  
2477 TGAACCTTGGAGAAATAAATCAGAACATGAGAAAGAGCTGGACTTGCATATAGGCT 2536 Db  
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2537 AATTTCTGGA-----GTAATAAACACATTAATTTGAATATCATATATCTATCAGATA 2589 Db  
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4641 AGAGTAAAAATATCAGTCATGGAATTAATATAGTGTCAAGAAAGTATGAGATGGAACCC 4700 Qy  
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4701 TTTCTTACTTTTTTACCTTCA-----TTTCTTAGTTTTTTTTTTTCTTTCACACCTGA 4752 Qy  
2709 TTTTTTTCTCTCTCTCCCATCAAGACCTTCCATTTAGTTTCTTCTTCTTCTTCTTCTCA 2768 Db  
4753 TCAGGCACTAGTAGACACCTATCTCTGTGAGCTATATATGACTTTTACAGCAACAAC 4812 Qy  
2769 ACAATCTCTTGGGAGCATTATCCATGTTGGGCTGGTGTACATTTCTATAGTAATGAT 2828 Db  
4813 ATTGCTGTGGCTCTTTGGGGAAGGAAACAGGATAGCAGGAGGCTCAGGCTAGCAAGT 4872 Qy  
2829 ACCATCATGTGGCTATTTTGGTGAAGAACA--ACAATGGAAGGCTTAGACTAACATA 2886 Db  
4873 CTGACTTGGCTTAAAGCCAGAGCATGTTGTATAGCAGAGAAAGTGAAGGCTTCTTCGCAAG 4932 Qy  
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4933 TGGGTGTGCTTAAGTAATCAGAAACAGGAGGCTCCGGTTGATGGAATATCATAGTAAAG- 4991 Qy  
2946 CAGGTACAATAAATCTCAGAAACATGAAGGCTCCAGTTGATGGAATTTTTCAGTAACNA 3005 Db  
4992 -----TATCTACCTTATCTCTCTATCGAACCTAAATCGTCTCTTTTCTTGTG 5042 Qy  
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5043 TGTAGGCTGATAAACACACTTGT--TTCTTTTGAAGTGTTCATGGCTTTGTAGATTTTA 5100 Qy  
3066 CATCATTTAATGAGTGTGCTGTTCTTCTTTGATAATTTGAAGGCTTTCTAGTTTTAA 3125 Db  
5101 GTGCTCTGGCAGTCTTGT--TAGAGGTTTGTACCTTGACACCTGGGCTTGGATGTTA 5158 Qy  
3126 TTGTGAAGCCAGTCTCTTGTATAGAACTATATCTAGACATGAGGAGGCTGAATGTTA 3185 Db  
5159 GCATGCCAAGGACACACTTCTGATGCTGTGTAAGGTTTATTTACTTACTTACT-- 5215 Qy  
3186 GCATGCCACAGACAGGCAATGCTTTTACATCTTGTCTTAAATAATTTACTGATTTCACTT 3245 Db  
5216 -----TTGTCTTTGGAAGGTGAAGGCTGTGAGAAAGAACTCACAGGAGATGTGTCT 5270 Qy  
3246 GCTTGTGTCTTTAGAAAAGTGAAGTGTGAGAGAGGAGATCTCATGTTCA----- 3296 Db  
5271 CTGTAGAAAACCTTTTTTTTCCCTTAAATGCTAATATCTTCACTTCACTCA--ACTT 5327 Qy  
3297 -----TCTGTGTGATTTTCAAGACCTTTAATCTCAATTTTGAAGAATCAAT 3342 Db  
5328 TGACTTTTATACCATCTCTGTCATGAAAGAGTGTGTAGCCCGCTCTCATGCTCTGGG 5387 Qy  
3343 TCATATTTGCAATGGGTGCTGATGGAAGAGTGAATTAATGCTTTTGTGCTGGTAGCTTCA 3402 Db





QY 6655 GCCTTCTAAAGAGCAATAGATCCCTGATCGACTTTT---ACTAAAGGAAGTG 6710  
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QY 6711 AGAAGCTAAAGCTCATCATATAGAGATTTCACATGAACCTGGCTCAGTTGAAAG 6770  
Db 696 GGAAGCCAACTCATCATGATGGGTGATTCGAATGAACCCCTGGTTAGTTACAAAG 755  
QY 6771 AAAATAGTGCAA--GTGTCCATAGAGACCAG--AGGTAGACTTGATTAACCAAGATTC 6827  
Db 756 GAAACCAATGCCACTTTTGTGTTATAAGACCAAGAGGTAGACTTCTTAAGCATAGATATT 815  
QY 6828 ATTGACAAATATTTTATGTCACATG---ATACACAGAAATAATGACTTTTAAA 6883  
Db 816 ATTGATAACATTTTCAITGTAACCTGGTGTCTATACAGAAACAATTTATTTTAAAT 875  
QY 6884 AATTGTTT-----GAAAGGAGGTACTCTCATCTCTTTA---GAAAAAAGCTTATG 6933  
Db 876 AATTGCTTTTTCATAAAAGATTAATTCTTCCATTCCTTTAGGGGAAAAACCCCTAAA 935  
QY 6934 TAACTTCA--TTTCCATATCCAAATATTTATATATGTAAGTTTATTTATTATAGTATA- 6990  
Db 936 TAGCTTCATGTTTCCATAATCAGTACTTTATATTTATAAATGTTATTTATTATTATAA 995  
QY 6991 -----CAITTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAACATTCG 7045  
Db 996 GACTGCATTTTATTTATATATCATTTTATTAATATGATTTATTTATAGAAACATTCGA 1055  
QY 7046 TATTGATA-TTTAGTATAAGCAATA--ATATTTATGACAAATCATG---AAAC 7097  
Db 1056 TATTGCTACTTGATGTAAGCTAAATGATGATTTATGACAAATATATAGACTATA 1115  
QY 7098 AAGATATCTTAGGCTTTTAAATAACACATGGATATC 7132  
Db 1116 CATGTTTATTTGACCTCAATAAACACTTGGATATC 1150

RESULT 7  
US-11-177-987-25  
; Sequence 25, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 25  
; LENGTH: 690  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-11-177-987-25  
Query Match 1.7%; Score 126; DB 7; Length 690;  
Best Local Similarity 71.7%; Pred. No. 7.1e-21;  
Matches 165; Conservative 0; Mismatches 65; Indels 0; Gaps 0;  
QY 2034 CTCCTCTCACTTATCAACTGTTGACACACTTGTGCGATCTCTGATGGCTGCTCAGAA 2093  
Db 29 CTCCTTCCCAGTCACAGTTGCTCGAGTTAGAAATGTTGCAATGTCGCGCCTGCAGAA 88

QY 2094 ATCTATGAGTTTTTCCCTTATGGGACTTTTGGCGCGCAGCTGCTTCTCTCATTTGCCCT 2153  
Db 89 ATCTGAGAGCTCTTTCTTATGGGACCTTGGCCACGAGTGGCTCTCTTCTTTGGCCT 148  
QY 2154 GTGGGCCAGGAGGCAAAATGCGTCCCGTCAACACCCGGTGCAAGCTTGAGGTGTC 2213  
Db 149 CTTGGTACAGGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGCGAGGCTTGACAAGTCAA 208  
QY 2214 CTTCCAGCAGCCCTACATCGTCAACCGCACCTTTATGCTGGCCCAAGAGG 2263  
Db 209 CTTCCAGCAGCCCTATATATCACAACCGCACCTTTCATGCTGGCTAAGGAGG 258  
RESULT 8  
US-11-177-987-18  
; Sequence 18, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 18  
; LENGTH: 418  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-11-177-987-18  
Query Match 1.6%; Score 122; DB 7; Length 418;  
Best Local Similarity 71.7%; Pred. No. 4.9e-20;  
Matches 175; Conservative 0; Mismatches 65; Indels 4; Gaps 1;  
QY 6535 ATAGCTTGGAGAGAGTGGAGAGATCAAGGGGATTTGGGAACTGGACCTGCTTTATGTC 6594  
Db 157 AAAGCTTGGAGAGAGTGGAGAGATCAAGGCAATTTGGAGAACTGGATTGCTGTTTATGTC 216  
QY 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAACGAGAACTGCTCTCTTCT 6654  
Db 217 TCTGAGAAATGCTTGGCAATTTGACCAGAGCAAGCTGAAAATGAATTAACCCCTTT 276  
QY 6655 GCCTTCTAAAAGAACAAATAGATCCCTGATGACTTTTTT---ACTAAAGGAAGTG 6710  
Db 277 CCCTGCTAGAAATAACAAATAGATGCCCAAGCGATTTTTTAAACCAAGGAGATG 336  
QY 6711 AGAAGCTAACGTCATCATCATATTAGAGATTTCATGAAACCTGGCTCAGTTGAAAAAG 6770  
Db 337 GGAAGCCAAACTCCATCATGATGGTGGATTCCAAATGAACCCCTGCGTTAGTTACAAAG 396  
QY 6771 AAAA 6774  
Db 397 GAAA 400

RESULT 9  
US-10-995-561-13277/c  
; Sequence 13277, Application US/10995561  
; Publication No. US20050272054A1  
; GENERAL INFORMATION:  
; APPLICANT: CARGILL, Michele et al.

; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH  
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF  
; TITLE OF INVENTION: DETECTION AND USES THEREOF  
; FILE REFERENCE: CL001559  
; CURRENT APPLICATION NUMBER: US/10/995,561  
; CURRENT FILING DATE: 2004-11-24  
; NUMBER OF SEQ ID NOS: 85702  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 13277  
; LENGTH: 173995  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; US-10-995-561-13277

Query Match 0.8%; Score 70; DB 6; Length 173995;  
Best Local Similarity 57.7%; Pred. No. 1.1e-05;  
Matches 164; Conservative 0; Mismatches 115; Indels 5; Gaps 2;  
QY 20 TTCTTCTAAATTAATAAAACCTATTTCTTAAATGAAAGCAACCCAGAGCGCTATTT 79  
DB 122980 TTATGTAATTTTATTTGTAATTTAGAAATAAATTTACAAAGTAAATAATTAATATATTT 122921  
QY 80 ATAGCAT- ---GGTGTCTGCACCATGCAGGTACAGATGGAATGTAAGGCGCTATTA 135  
DB 122920 ATGGTGATTAATCTGTTTGGATATAGTATGATGATGTAAGTGGCTAAAGATGTTTAAACA 122861  
QY 136 TCAGCATTAACCAACATGTTTAAATGTTTCTTCTGCAAGCAAACTTGAATCTATGCTTT 195  
DB 122860 CAGCATTAACCTACATAGTCTTTTCTTGTGATGAACATTAATAATGATTTCT 122801  
QY 196 AACAATCTTCAAGCCTCTAATATAGTCTAAGCAAGTGGAGTCCGC-TGCTGTCCAAACAG 254  
DB 122800 TAACAATTTTCAATAATTTCAATATATGTTTAACTGTAATCATCATGATGTACACAG 122741  
QY 255 AGCTCTTGAGCAGCTCTCTCTCTCTTTGCAATTTTATGTTCTTT 298  
DB 122740 ATCTCTCAAACTTATTTTCTTCTAACTGAAATTTTGTGTCATTT 122697

RESULT 10  
US-10-995-561-13314/c  
; Sequence 13314, Application US/10995561  
; Publication No. US2005027054A1  
; GENERAL INFORMATION:  
; APPLICANT: CARGILL, Michele et al.  
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH  
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF  
; TITLE OF INVENTION: DETECTION AND USES THEREOF  
; FILE REFERENCE: CL001559  
; CURRENT APPLICATION NUMBER: US/10/995,561  
; CURRENT FILING DATE: 2004-11-24  
; NUMBER OF SEQ ID NOS: 85702  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 13314  
; LENGTH: 119036  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: misc feature  
; LOCATION: (1)...(119036)  
; OTHER INFORMATION: n = A,T,C or G, or insertion/deletion polymorphism (see Tables 1-13314)

Query Match 0.8%; Score 62.6; DB 6; Length 119036;  
Best Local Similarity 47.9%; Pred. No. 0.00057;  
Matches 241; Conservative 0; Mismatches 259; Indels 3; Gaps 2;  
QY 6871 ATGTACTTTTAAAAAATTTGTTGAAAGGAGGTACTCTCATCTTTAGAAAAAAGCTT 6930  
DB 31607 ATATATTTTATATTTTATATATAAATCTTATTTTATATATATTTTATATATAAAAA 31548  
QY 6931 ATGTAATCTCATTTCCATATCCAAATATTTTATATATGTAAGTTTATTTATTAAGTATA 6990

DB 31547 ATCATATTTTATATATATATTTTATATATATTTTATATATATATAAATCTTTATTTATATAT-TA 31489  
QY 6991 CATTTATTTATGTCAGTCTTTTATTAATATGGAATTTATTTATAGAAACATATCTGCTATTG 7050  
DB 31488 TTTTATATATATTTTATATATATATAAATCTTATTTTATATATATTTTATATATATATA 31429  
QY 7051 ATATTTAGTATAAGGCAATAATATTTTATGACATACTATGGAACAACAAGATATCTTAGG 7110  
DB 31428 AAATCTTATTTTATATATATTTTATATATATAAATCTTATTTTATATATATTTTATATA 31369  
QY 7111 CTTTAAATAACACATGATATCAATCT--TCTGCTTGTAAATTTTCTCCCTTTAAT 7168  
DB 31368 TATATATAAATCTTATTTTATATATATTTTATATATATAAATCTTATTTTATATAT 31309  
QY 7169 ATCAACAATACCATCATCATCATCATCATCATCATCATCATCATCATCATCATCATCATCAT 7228  
DB 31308 ATTTTATATATATAAATCTTATTTTATACAGCCATTTTCTAATAAATAAATAACAGCC 31249  
QY 7229 ATATTAATCTGTTAAAGTTGGTTCCTGGAGCCCTGCGTTTTTGTGTGTGTGTGTGTGTG 7288  
DB 31248 ATTTTCTAAATATAAATTTTCTTTTAAACTTTTTCCTTTTCCAGGGGCAATGC 31189  
QY 7289 TGTGGGGTTATGATGTGAAGCCAGAGATGATATAGTGTCTCTCTATCAGTCTT 7348  
DB 31188 TCACTGATATAAATAATGATGTTATAAATATTTTCTTTTCTTTTCTTTTCTTTTCTTTT 31129  
QY 7349 TGCCTTATTTTGGAGACAGGCT 7371  
DB 31128 AGTCTGCTTTGTCACCCAGGCT 31106

RESULT 11  
US-11-140-417-22  
; Sequence 22, Application US/11140417  
; Publication No. US20050266515A1  
; GENERAL INFORMATION:  
; APPLICANT: O'Brien, Deborah A  
; TITLE OF INVENTION: GLYCERALDEHYDE 3-PHOSPHATE DEHYDROGENASE-S (GAPDS), A GLYCOLYTIC  
; TITLE OF INVENTION: ENZYME EXPRESSED ONLY IN MALE GERM CELLS, IS A TARGET FOR MALE  
; TITLE OF INVENTION: CONTRACEPTION  
; FILE REFERENCE: 421/76/2 PCT/CIP  
; CURRENT APPLICATION NUMBER: US/11/140,417  
; CURRENT FILING DATE: 2005-05-27  
; PRIOR APPLICATION NUMBER: US 60/429,638  
; PRIOR FILING DATE: 2002-11-27  
; PRIOR APPLICATION NUMBER: PCT/US2003/037800  
; PRIOR FILING DATE: 2003-11-26  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 22  
; LENGTH: 11462  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; US-11-140-417-22

Query Match 0.8%; Score 61.6; DB 7; Length 11462;  
Best Local Similarity 63.1%; Pred. No. 0.00023;  
Matches 111; Conservative 0; Mismatches 64; Indels 1; Gaps 1;  
QY 434 TGTATGTGTGTGCACCAAGTGTGCTGCTGTGTTGTGGGGCAAGGAGCAGG-AGAGGGT 492  
DB 27 TATGTATGTGTGAACCTGTGCGTACGGGTGCTGCAGAGGCGCAGAGAGGCGCATCAGAT 86  
QY 493 GCCTTGGCAGGAGTACGGATGTTGTGAGCCACCATGAGGATGCTGGAGTTAGACC 552  
DB 87 CCTCTGGAAGTGGAGTTACCAATGTTGGAATTTCTACGTGGGTGCTGGGATCTGAACT 146  
QY 553 CAGGTCTCTCCAGAGTGCAGCAATGCTCTTAAACCAACACAGCAGCATTTCTCTCTC 608  
DB 147 CGAGTCTCTGTGAGAGCAACAATGCTCTTAACTACTGAGTACTTCTCTGTCCTC 202

## RESULT 12

US-10-933-025-24/c  
; Sequence 24, Application US/10933025  
; Publication No. US20050265987A1  
; GENERAL INFORMATION:  
; APPLICANT: ROSEN, STEVEN  
; APPLICANT: HEMMERICH, STEFAN  
; APPLICANT: TOMITA, MEGUMI  
; TITLE OF INVENTION: Sulfotransferases and methods of use  
; FILE REFERENCE: USAL-230CON  
; CURRENT APPLICATION NUMBER: US/10/933,025  
; CURRENT FILING DATE: 2004-09-01  
; PRIOR APPLICATION NUMBER: 10/025,966  
; PRIOR FILING DATE: 2001-12-21  
; PRIOR APPLICATION NUMBER: 60/258,577  
; PRIOR FILING DATE: 2000-12-27  
; PRIOR APPLICATION NUMBER: 60/267,831  
; PRIOR FILING DATE: 2001-09-02  
; NUMBER OF SEQ ID NOS: 26  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 24  
; LENGTH: 171936  
; TYPE: DNA  
; ORGANISM: Mus musculus  
US-10-933-025-24

Query Match 0.8%; Score 59.2; DB 6; Length 171936;  
Best Local Similarity 83.8%; Pred. No. 0.0049;  
Matches 67; Conservative 0; Mismatches 13; Indels 0; Gaps 0;

Qy 508 TCACGGATGGTGTGAGCCACCATGAGGATCTGGGAGTTAGACCCAGGTCCTCCAGAAG 567  
Db 71888 TTACGGGTGGTGTGAGCCACCATGTTGCTGGGATTTGAACCTCAGGACCTCCAGAAG 71829

Qy 568 TGCAAGCAATGCTCTTAACC 587  
Db 71828 AGCAGTCAGTCTCTTAACC 71809

RESULT 13  
US-11-121-086-105  
; Sequence 105, Application US/11121086  
; Publication No. US20050266459A1  
; GENERAL INFORMATION:  
; APPLICANT: POULSEN, TIM S.  
; APPLICANT: NIELSEN, KIRSTEN V.  
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES  
; FILE REFERENCE: 09138.6000-00000  
; CURRENT APPLICATION NUMBER: US/11/121,086  
; CURRENT FILING DATE: 2005-05-04  
; PRIOR APPLICATION NUMBER: 60/567,570  
; PRIOR FILING DATE: 2004-05-04  
; NUMBER OF SEQ ID NOS: 107  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 105  
; LENGTH: 171486  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-11-121-086-105

Query Match 0.8%; Score 58.4; DB 7; Length 171486;  
Best Local Similarity 49.5%; Pred. No. 0.0077;  
Matches 206; Conservative 0; Mismatches 206; Indels 4; Gaps 2;

Qy 6826 TCATTGCAATATTTTATGTGCTGATGATACACAGAAAAATAATGCTACTTTAAAAAA 6885  
Db 16176 TTATATAGATATAATATTTATGTTATATATATATATATATATATATATATATATATA 16235

Qy 6886 TTGTTGAAAGGAGGTTACCTCACTTCCTTTAGAAAAAGGTTATGTAACCTTCATTTTC 6945  
Db 16236 ATTATTAAATATGTTTATATATAAATTAATATATATATATATATATATATATATATAT 16292

## RESULT 14

US-10-995-561-13460  
; Sequence 13460, Application US/10995561  
; Publication No. US20050272054A1  
; GENERAL INFORMATION:  
; APPLICANT: CARGILL, Michele et al.  
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH  
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF  
; TITLE OF INVENTION: DETECTION AND USES THEREOF  
; FILE REFERENCE: CL001559  
; CURRENT APPLICATION NUMBER: US/10/995,561  
; CURRENT FILING DATE: 2004-11-24  
; NUMBER OF SEQ ID NOS: 85702  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 13460  
; LENGTH: 20317  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-10-995-561-13460

Query Match 0.8%; Score 57.4; DB 6; Length 20317;  
Best Local Similarity 46.3%; Pred. No. 0.0035;  
Matches 244; Conservative 5; Mismatches 271; Indels 7; Gaps 2;

Qy 6725 ATCATCATTTAGAAGATTTTCACATGCAACCTGGCTGCTGAAAAAGAAAAATAGTGTCAAG 6784  
Db 18098 ATATAYWATATATGATATACATCTATATATAATATAATATATATATATATATATATAT 18157

Qy 6785 TTGTCCATGAGACGACAGAGGTAGACTTCGATAACCAAGAGATTCATTGACCAATATTTTATT 6844  
Db 18158 TAATATATGATATATGAT 18217

Qy 6845 GTCACCTGATGATACACAGAAAAATAATGCTCTTTAAAAAATGTTTGAAGAGGTTTAC 6904  
Db 18218 ATATCAT 18277

Qy 6905 CTCCTCATTCCTTTAGAAAAAGGCTTATGTAACCTTCATTTCCTATCCCAATATTTTATAT 6964  
Db 18278 ATGATATATCATATATCATATATAATATATATATATATATATATATATATATATATATAT 18337

Qy 6965 ATGTAAAGTTTATTTATTAAGTATACATATTTTAT -TTATGTGCTGCTTTTATTAATATGGATT 7023  
Db 18338 ATATTTTATATATTTAT 18397

Qy 7024 TATTTATAGAAACATTTATCTGCTATTTGATATTTTAGTATAAGGCAAAATAATTTATGACA 7083  
Db 18398 TATATATATATAATAATTTTACATATAATAATAATAATAATAATAATAATAATAATAATA 18457

Qy 7084 ATAACATA-----TGGAAACAAGATCTTAGGCTTTTAAATAAACAACATGGGATATCAATA 7137  
Db 18458 TAATATAATAATAGGATATATAATAATAATAATAATAATAATAATAATAATAATAATAATA 18517

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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 125.756 Seconds  
(without alignments)  
15704.028 Million cell updates/sec

Title: US-09-751-797-9

Perfect score: 1111

Sequence: 1 aacaggctctctcagtt.....aaacacatgaatatcataaa 1111

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 1303057 seqs, 888780828 residues

Total number of hits satisfying chosen parameters: 2606114

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Issued Patents NA.\*

- 1: /cgn2\_6/ptodata/1/ina/1 COMB.seq.\*
- 2: /cgn2\_6/ptodata/1/ina/5 COMB.seq.\*
- 3: /cgn2\_6/ptodata/1/ina/6A COMB.seq.\*
- 4: /cgn2\_6/ptodata/1/ina/6B COMB.seq.\*
- 5: /cgn2\_6/ptodata/1/ina/H COMB.seq.\*
- 6: /cgn2\_6/ptodata/1/ina/PCPUS COMB.seq.\*
- 7: /cgn2\_6/ptodata/1/ina/PP COMB.seq.\*
- 8: /cgn2\_6/ptodata/1/ina/RE COMB.seq.\*
- 9: /cgn2\_6/ptodata/1/ina/backfiles1.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1111	100.0	1111	3	US-09-178-973B-9
2	1111	100.0	1111	3	US-09-419-568F-9
3	1111	100.0	1111	3	US-09-354-243B-9
4	1049.4	94.5	1166	3	US-10-084-298-3
5	1048.4	94.4	1050	3	US-10-090-365-40
6	1047.8	94.3	1119	3	US-09-178-973B-7
7	1047.8	94.3	1119	3	US-09-419-568F-7
8	1047.8	94.3	1119	3	US-09-354-243B-7
9	602.4	54.2	5935	3	US-09-178-973B-17
10	602.4	54.2	5935	3	US-09-419-568F-29
11	602.4	54.2	5935	3	US-09-354-243B-29
12	555.2	50.0	7445	3	US-09-178-973B-8
13	555.2	50.0	7445	3	US-09-419-568F-8
14	555.2	50.0	7445	3	US-09-354-243B-8
15	529	47.6	1191	3	US-10-084-298-1
16	525.6	47.3	1152	3	US-09-870-574-1
17	525.2	47.3	1116	3	US-10-090-365-14
18	525.2	47.3	1116	3	US-09-728-911-14
19	407.6	36.7	689	3	US-09-949-016-5443
20	407.6	36.7	690	3	US-09-419-568F-24
21	407.6	36.7	690	3	US-09-354-243B-24
22	217.4	19.6	8888	3	US-09-949-016-17185
23	185.2	16.7	191	3	US-10-084-298-9
24	127.6	11.5	4797	3	US-09-419-568F-25

ALIGNMENTS

RESULT 1

US-09-178-973B-9

; Sequence 9, Application US/09178973B

; Patent No. 6274710

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Louhed, Jamila

; APPLICANT: Renauld, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; TITLE OF INVENTION: (TIPS)

; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5543

; CURRENT APPLICATION NUMBER: US/09/178,973B

; CURRENT FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 17

; SEQ ID NO 9

; LENGTH: 1111

; TYPE: DNA

; ORGANISM: Mus musculus

US-09-178-973B-9

Query Match 100.0%; Score 1111; DB 3; Length 1111;

Best Local Similarity 100.0%; Pred. No. 3.4e-274;

Matches 1111; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGCGGATCGTGATGGCTGTCTCT	60
Db	1	AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGCGGATCGTGATGGCTGTCTCT	60
Qy	61	GCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTGGCCGCCAGCTGCCTGTCTCAT	120
Db	61	GCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTGGCCGCCAGCTGCCTGTCTCAT	120
Qy	121	TGCCCTGTGGGCCAGAGGCAAAATGGCTGCCCATCAACCCGGTGCAAGCTTGAGGT	180
Db	121	TGCCCTGTGGGCCAGAGGCAAAATGGCTGCCCATCAACCCGGTGCAAGCTTGAGGT	180
Qy	181	GTCCAACTTCCACGACCGGTACATCGTCAACCCGACCTTTATGCTGCGCAAGAGGCCAG	240
Db	181	GTCCAACTTCCACGACCGGTACATCGTCAACCCGACCTTTATGCTGCGCAAGAGGCCAG	240
Qy	241	CCTTGAGATAAACAACAACAGACGCTCGGGCTCATCGGGGAGAAACTGTTCGAGAGTCAG	300
Db	241	CCTTGAGATAAACAACAACAGACGCTCGGGCTCATCGGGGAGAAACTGTTCGAGAGTCAG	300
Qy	301	TGCTAAGGATCAGTGTCTGATGAGGAGGCTCAACTTCACTCCCTGGAAGACATCTCT	360
Db	301	TGCTAAGGATCAGTGTCTGATGAGGAGGCTCAACTTCACTCCCTGGAAGACATCTCT	360



Db 901 CCTATGTAACCTCATTCCATAACCAATACCTTTATATATATGTAAGTTTATTTATATAAGT 960  
Qy 961 ATACATTTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAAAATTTATCTGATG 1020  
Db 961 ATACATTTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAAAATTTATCTGATG 1020  
Qy 1021 TTGATATTTGAGTATAAAGCAATAATATTTATTTATGATTAATACTATAGAAAACAAGATATCT 1080  
Db 1021 TTGATATTTGAGTATAAAGCAATAATATTTATTTATGATAATACTATAGAAAACAAGATATCT 1080  
Qy 1081 TAGGCTTTTATTAACACATCAATATCATATA 1111  
Db 1081 TAGGCTTTTATTAACACATCAATATCATATA 1111

RESULT 3  
US-09-354-243B-9  
; Sequence 9, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Pa  
; TITLE OF INVENTION: (TIFs)  
; FILE REFERENCE: LUD 5543.1  
; CURRENT APPLICATION NUMBER: US/09/354,243B  
; CURRENT FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-354-243B-9

Query Match 100.0%; Score 1111; DB 3; Length 1111;  
Best Local Similarity 100.0%; Pred. No. 3.4e-274; Mismatches 0; Indels 0; Gaps 0;  
Matches 1111; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AACAGGCTCTCCTCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGCTGTCT 60  
Db 1 AACAGGCTCTCCTCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGCTGTCT 60  
Qy 61 GCAGAAATCTATGAGTTTCCCTTATGCGGACTTTTGGCGCCGAGCTGCTTCTCAT 120  
Db 61 GCAGAAATCTATGAGTTTCCCTTATGCGGACTTTTGGCGCCGAGCTGCTTCTCAT 120  
Qy 121 TGCCTGTGGGCCAGAGGCAATGCGTCCCATCAACACCCGGTGCAAGCTTGAGGT 180  
Db 121 TGCCTGTGGGCCAGAGGCAATGCGTCCCATCAACACCCGGTGCAAGCTTGAGGT 180  
Qy 181 GTCCAACTTCCAGAGCGGTACATCGTCAACCCGACCTTTATGCTGGCAAGGAGCCAG 240  
Db 181 GTCCAACTTCCAGAGCGGTACATCGTCAACCCGACCTTTATGCTGGCAAGGAGCCAG 240  
Qy 241 CCTTGAGATPAAACAACAGAGCTCCGGCTCATCGGGGAGAAACTGTTCGAGAGGTGAG 300  
Db 241 CCTTGAGATPAAACAACAGAGCTCCGGCTCATCGGGGAGAAACTGTTCGAGAGGTGAG 300  
Qy 301 TGCTAAGGATCATGCTTACTGATGAGAGGCTCACTTCCCTGGAAGACATTTCT 360  
Db 301 TGCTAAGGATCATGCTTACTGATGAGAGGCTCACTTCCCTGGAAGACATTTCT 360  
Qy 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATGACGAGGTGGTGTCTTCTGACCAA 420  
Db 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATGACGAGGTGGTGTCTTCTGACCAA 420  
Qy 421 ACTCAGCAATCAGCTCAGTCTCTGTCAATCAGTGTGACGACCAAGAAATCCAGAAAG 480

Db 421 ACTCAGCAATCAGCTCAGCTCCTGTCCATCATCAGTGTGACGACCAAGAAATCCAGAAAG 480  
Qy 481 TGTCAAGAGGCTCAAGAGAGACAGTGAAGAGCTTTGGAGAGAGCGGAGAGATCAAGCGAT 540  
Db 481 TGTCAAGAGGCTCAAGAGAGACAGTGAAGAGCTTTGGAGAGAGCGGAGAGATCAAGCGAT 540  
Qy 541 CGGGAACTGGACCTGCTGTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAAGAAAG 600  
Db 541 CGGGAACTGGACCTGCTGTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAAGAAAG 600  
Qy 601 CTAGAAAAAGAGAGAGCTGCTTCTGCTTCTTAAAGAAACAATAAGATCCCTGATG 660  
Db 601 CTAGAAAAAGAGAGAGCTGCTTCTGCTTCTTAAAGAAACAATAAGATCCCTGATG 660  
Qy 661 GACTTTTTTACTAAAGAAAGTGAGAGAGCTTAAGCTCCACCATCATTTAGAGATTTTCACAT 720  
Db 661 GACTTTTTTACTAAAGAAAGTGAGAGAGCTTAAGCTCCACCATCATTTAGAGATTTTCACAT 720  
Qy 721 GAAACCTGCTCAGTTGAAAGAGAAATAGTGTCAAGTTGTCCATGAGACCGAGGTAGA 780  
Db 721 GAAACCTGCTCAGTTGAAAGAGAAATAGTGTCAAGTTGTCCATGAGACCGAGGTAGA 780  
Qy 781 CTTGATAACCAACAGAGATTCATTGACAATATTTTATTTGCTCATGATTAATGCAACAGAAA 840  
Db 781 CTTGATAACCAACAGAGATTCATTGACAATATTTTATTTGCTCATGATTAATGCAACAGAAA 840  
Qy 841 AGTATGCTACTTTAAAAAATTTGTTGAAAGAGGTTACCTCTCATTTCTTAGAAGAAAG 900  
Db 841 AGTATGCTACTTTAAAAAATTTGTTGAAAGAGGTTACCTCTCATTTCTTAGAAGAAAG 900  
Qy 901 CCTATGTAACCTCATTTCCATAACCAATCTTTTATATATGTAAGTTTATTTATTAAGT 960  
Db 901 CCTATGTAACCTCATTTCCATAACCAATCTTTTATATATGTAAGTTTATTTATTAAGT 960  
Qy 961 ATACATTTTATTTATGCTCAGTTTATTAATATGATTTATTTATAGAAATTTATCTGATG 1020  
Db 961 ATACATTTTATTTATGCTCAGTTTATTAATATGATTTATTTATAGAAATTTATCTGATG 1020  
Qy 1021 TTGATATTTTGGATATAAGCAATAATATTTATGATAATACTATAGAAAACAAGATATCT 1080  
Db 1021 TTGATATTTTGGATATAAGCAATAATATTTATGATAATACTATAGAAAACAAGATATCT 1080  
Qy 1081 TAGGCTTTTATTAACACATCAATATCATATA 1111  
Db 1081 TAGGCTTTTATTAACACATCAATATCATATA 1111

RESULT 4  
US-10-084-298-3  
; Sequence 3, Application US/10084298  
; Patent No. 6939545  
; GENERAL INFORMATION:  
; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: Xuan, Dejun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory  
; TITLE OF INVENTION: Disorders  
; FILE REFERENCE: G15358 CIP  
; CURRENT APPLICATION NUMBER: US/10/084,298  
; CURRENT FILING DATE: 2002-09-10  
; PRIOR APPLICATION NUMBER: 60/270,823  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR APPLICATION NUMBER: 60/281,353  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR APPLICATION NUMBER: 60/131,473  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR APPLICATION NUMBER: 09/561,811  
; PRIOR FILING DATE: 2000-04-28  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 3

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; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Murine
US-10-084-298-3

Query Match      94.5%; Score 1049.4; DB 3; Length 1166;
Best Local Similarity 97.1%; Pred. No. 2e-258;
Matches 1079; Conservative 0; Mismatches 31; Indels 1; Gaps 1;

Qy 1 AACAGGCTCTCCTCTCAGTTATCAACTTTTGACACTTTGTGGGATCTCTGATGGCTGTCT 60
Db 26 AACAGGCTCTCCTCTCAGTTATCAACTTTTGACACTTTGTGGGATCTCTGATGGCTGTCT 85

Qy 61 GCAGAAATCTATGATGTTTTCCCTTTATGGGACTTTTGGCCGCGCAGCTGCCTTCTTCAT 120
Db 86 GCAGAAATCTATGATGTTTTCCCTTTATGGGACTTTTGGCCGCGCAGCTGCCTTCTTCAT 145

Qy 121 TGGCCTGTGGGCCCCAGGAGGCAATGCGCTGCCCATCAACACCCGGTGCAAGCTTGAGGT 180
Db 146 TGGCCTGTGGGCCCCAGGAGGCAATGCGCTGCCCATCAACACCCGGTGCAAGCTTGAGGT 205

Qy 181 GTCCAACTTCAGAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGAGGCCAG 240
Db 206 GTCCAACTTCAGAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGAGGCCAG 265

Qy 241 CCTTGCAGATAACAACACAGAGCTCCGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 300
Db 266 CCTTGCAGATAACAACACAGAGTCCGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 325

Qy 301 TGCTAAGGATCAGTGTCTACCTGTATGAAGCAGGTGCTCAACTTTCACCCCTGGAAGCATTC 360
Db 326 TGCTAAGGATCAGTGTCTACCTGTATGAAGCAGGTGCTCAACTTTCACCCCTGGAAGCATTC 385

Qy 361 GCTCCCCAGTCAAGAGGTTCCGGCCCTCATGTCAGAGAGTGTGCTTCTTCGACCA 420
Db 386 GCTCCCCAGTCAAGAGGTTCCAGCCCTCATGTCAGAGAGTGTGCTTCTTCGACCA 445

Qy 421 ACTCAGCAATCAGCTCAGCTCCTGTCAATCAGTGTGAGCAGCAGTGTGAGTCCAGAGAA 480
Db 446 ACTCAGCAATCAGCTCAGCTCCTGTCAATCAGCAGGTTGAGCAGCAGTGTGAGTCCAGAGAA 505

Qy 481 TGTCAAGAGGCTGAAGGAGACAGTGAAGAAAGCTTTGGAGAGCGGAGAGATCAAGCGAT 540
Db 506 TGTCAAGAGGCTGAAGGAGACAGTGAAGAAAGCTTTGGAGAGTGTGAGAGATCAAGCGAT 565

Qy 541 CGGGGAACTGGAACCTGTGTTTATGCTCTGAGAAATGCTTGTGCTGAGCGAGAGAAAG 600
Db 566 TGGGGAACTGGAACCTGTGTTTATGCTCTGAGAAATGCTTGTGCTGAGCGAGAGAAAG 625

Qy 601 CTAGAAACGAAAGTCACTCTCCCTGCTTCTTAAAGAAAGCAATTAAGATCCCTGAAATG 660
Db 626 CTAGAAACGAAAGTCACTCTCCCTGCTTCTTAAAGAAAGCAATTAAGATCCCTGAAATG 685

Qy 661 GACTTTTTTACTAAGGAAAGTGAGAAAGCTAACCGTCCACCATCAATTAGAAGATTTACAT 720
Db 686 GACTTTTTTACTAAGGAAAGTGAGAAAGCTAACCGTCCATCAATTAGAAGATTTACAT 745

Qy 721 GAAACCTGGCTCAGTTGAAGAGAAATAGTGTCAAGTGTGTCATGAGACAGAGGTAGA 780
Db 746 GAAACCTGGCTCAGTTGAAGAGAAATAGTGTCAAGTGTGTCATGAGACAGAGGTAGA 805

Qy 781 CTTGATACCAACAAGATTCATTCGACATATTTTATGTCATGATATGATCAACAGAGAA 840
Db 806 CTTGATACCAACAAGATTCATTCGACATATTTTATGTCATGATATGATCAACAGAGAA 865

Qy 841 AGTATGTACTTTTAAAAAATTTGTTGAAAGGAGGTTACCTCTCATTTCTAGAGAGAAAG 900
Db 866 ATAATGTACTTTTAAAAAATTTGTTGAAAGGAGGTTACCTCTCATTTCTTTAGAGAAAG 925

Qy 901 CCTATGTAACCTTCATTTCCATAACCAATATCTTTATATATATGTAAGTATTTATTTAAGT 960
Db 926 CTTATGTAACCTTCATTTCCATAACCAATATTTTATATATATGTAAGTATTTATTTAAGT 985
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Qy 961 ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAAAATTTATCTGATG 1020
Db 986 ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAAAATTTATCTGCTA 1045

Qy 1021 TTGATATTTGAGTATAAAGCAATAATATTTATGATAATTAACACTATAGAAAACAAGATATCT 1080
Db 1046 TTGATATTT-AGTATAAGGCAATAATATTTATGACAATAACTATGGAACAAGATATCT 1104

Qy 1081 TAGGCTTTAATAAACACATGAATATCATAAA 1111
Db 1105 TAGGCTTTAATAAACACATGGATATCATAAA 1135

RESULT 5
US-10-090-365-40
; Sequence 40, Application US/10090365
; Patent No. 6875845
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Chen, Zhi
; TITLE OF INVENTION: Mouse Cytokine Receptor
; FILE REFERENCE: 01-08
; CURRENT APPLICATION NUMBER: US/10/090,365
; CURRENT FILING DATE: 2002-03-04
; PRIOR APPLICATION NUMBER: US 60/273,035
; PRIOR FILING DATE: 2001-03-02
; PRIOR APPLICATION NUMBER: US 60/279,232
; PRIOR FILING DATE: 2001-03-27
; NUMBER OF SEQ ID NOS: 49
; SOFTWARE: Fast-Seq for Windows Version 3.0
; SEQ ID NO 40
; LENGTH: 1050
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (50)...(589)
US-10-090-365-40

Query Match      94.4%; Score 1048.4; DB 3; Length 1050;
Best Local Similarity 99.9%; Pred. No. 3.5e-258;
Matches 1049; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 AACAGGCTCTCCTCTCAGTTATCAACTTTTGACACTTTGTGGGATCGGTGATGGCTGTCT 60
Db 1 AACAGGCTCTCCTCTCAGTTATCAACTTTTGACACTTTGTGGGATCGGTGATGGCTGTCT 60

Qy 61 GCAGAAATCTATGATGTTTTCCCTTTATGGGACTTTTGGCCGCGCAGCTGCCTTCTTCAT 120
Db 61 GCAGAAATCTATGATGTTTTCCCTTTATGGGACTTTTGGCCGCGCAGCTGCCTTCTTCAT 120

Qy 121 TGGCCTGTGGGCCCCAGGAGGCAAAATGCGCTGCCCATCAACACCCGGTGCAAGCTTGAGGT 180
Db 121 TGGCCTGTGGGCCCCAGGAGGCAAAATGCGCTGCCCATCAACACCCGGTGCAAGCTTGAGGT 180

Qy 181 GTCCAACTTCAGAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGAGGCCAG 240
Db 181 GTCCAACTTCAGAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGAGGCCAG 240

Qy 241 CTTTGCAGATAACAACACAGAGCTCCGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 300
Db 241 CTTTGCAGATAACAACACAGAGCTCCGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 300

Qy 301 TGCTAAGGATCAGTGTCTACCTGTATGAAGCAGGTGCTCAACTTTCACCTGGAAGCATTC 360
Db 301 TGCTAAGGATCAGTGTCTACCTGTATGAAGCAGGTGCTCAACTTTCACCTGGAAGCATTC 360

Qy 361 GCTCCCCAGTCAAGAGGTTCCGGCCCTCATGTCAGAGAGTGTGCTTCTTCGACCA 420
Db 361 GCTCCCCAGTCAAGAGGTTCCGGCCCTCATGTCAGAGAGTGTGCTTCTTCGACCA 420
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Qy	421	ACTCAGCAATCAGCTCAGCTCCTGTGCATCATCAGTGGTGA	CGACCGACAGCATCCAGAAGAA	480
Db	421	ACTCAGCAATCAGCTCAGCTCCTGTGCATCATCAGTGGTGA	CGACCGACAGCATCCAGAAGAA	480
Qy	481	TGTCAGAAAGGCTGAAGGAGACAGTGA	AAAAAGCTTTGGAGAGAGCGGAGAGATCA	540
Db	481	TGTCAGAAAGGCTGAAGGAGACAGTGA	AAAAAGCTTTGGAGAGAGCGGAGAGATCA	540
Qy	541	CGGGGAACCTGGAACCTGCTGTTTATGCTCTCGAGAAATGCTTGGCTCTGACGCGAGAAGAAG	600	
Db	541	CGGGGAACCTGGAACCTGCTGTTTATGCTCTCGAGAAATGCTTGGCTCTGACGCGAGAAGAAG	600	
Qy	601	CTAGAAAAACGAAGAACTGCTCCTTCCCTGCTCTTAAAAAGAAACAATAAGATCCCTGAATG	660	
Db	601	CTAGAAAAACGAAGAACTGCTCCTTCCCTGCTCTTAAAAAGAAACAATAAGATCCCTGAATG	660	
Qy	661	GACTTTTTTTACTTAAAGGAAAGTGAAGAAGCTAAAGCTCCACCATCATTTAGAGAAGATTTCCACAT	720	
Db	661	GACTTTTTTTACTTAAAGGAAAGTGAAGAAGCTAAAGCTCCACCATCATTTAGAGAAGATTTCCACAT	720	
Qy	721	GAACACCTGGCTCAGTTGAAAGAGAGAAAAATAGTGTCAAGTTGTCTCCATGAGACCAGAGGTAGA	780	
Db	721	GAACACCTGGCTCAGTTGAAAGAGAGAAAAATAGTGTCAAGTTGTCTCCATGAGACCAGAGGTAGA	780	
Qy	781	CTTGATTAACACAAAGATTCATTGACAAATATTTTATGTCAATGATTAATGCCAAGAAAA	840	
Db	781	CTTGATTAACACAAAGATTCATTGACAAATATTTTATGTCAATGATTAATGCCAAGAAAA	840	
Qy	841	AGTATGTACTTTTAAAAAATGTTTGAAGGAGGTTTACCTCTCATTCCTCTAGAGAAAG	900	
Db	841	AGTATGTACTTTTAAAAAATGTTTGAAGGAGGTTTACCTCTCATTCCTCTAGAGAAAG	900	
Qy	901	CCTATGTAACTTCATTTTCCATAACCAATACCTTTATATATGTAAAGTTTATTTATATAAGT	960	
Db	901	CCTATGTAACTTCATTTTCCATAACCAATACCTTTATATATGTAAAGTTTATTTATATAAGT	960	
Qy	961	ATACATTTTATTTAATGTCAGTTTATTAATATGGAATTTATTTATAGAAAAATATCTGATG	1020	
Db	961	ATACATTTTATTTAATGTCAGTTTATTAATATGGAATTTATTTATAGAAAAATATCTGATG	1020	
Qy	1021	TTGATATTTTCAGTATAAAGCAAAATATTT	1050	
Db	1021	TTGATATTTTCAGTATAAAGCAAAATATTT	1050	

RESULT 6

US-09-178-973B-7

; Sequence 7, Application US/09178973B

; Patent No. 6274710

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Louhed, Jamila

; APPLICANT: Renaud, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fc

; TITLE OF INVENTION: (TIPI)

; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5543

; CURRENT APPLICATION NUMBER: US/09/178,973B

; CURRENT FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 17

; SEQ ID NO 7

; LENGTH: 1119

; TYPE: DNA

; ORGANISM: Mus musculus

US-09-178-973B-7

Query Match	94.3%	Score 1047.8;	DB 3;	Length 1119;
Best Local Similarity	97.0%	Pred. No. 5.1e-258;		
Matches 1078;	Conservative 0;	Mismatches 32;	Indels 1;	Gaps 1;

Qy	1	AACAGGCTCTCCTCTCAGTTATCACTTTTGAACATTTGTGCGATCGGTGATGCTGTCT	60
Db	3	AACAGGCTCTCCTCTCAGTTATCACTTTTGAACATTTGTGCGATCTCTGATGCTGTCT	62

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RESULT 7
US-09-419-568F-7
; Sequence 7, Application US/09419568F
; Patent No. 6331613
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; FILE REFERENCE: LUD 5543.2
; CURRENT APPLICATION NUMBER: US/09/419,568F
; PRIOR FILING DATE: 1999-10-18
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 7
; LENGTH: 1119
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-09-419-568F-7

Query Match 94.3%; Score 1047.8; DB 3; Length 1119;
Best Local Similarity 97.0%; Pred. No. 5.1e-258;
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;

Qy 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCGGTGATGGCTGTCT 60
Db 3 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCTGTGATGGCTGTCT 62

Qy 61 GCAGAAATCTATAGTTTTTCCCTTATGGGACCTTTGGCCGCCAGCTGCTCTTCAT 120
Db 63 GCAGAAATCTATAGTTTTTCCCTTATGGGACCTTTGGCCGCCAGCTGCTCTTCAT 122

Qy 121 TGGCCTGTGGCCCGAGGAGCAATGCGTGCCTCAACACCCGGTGCAAGCTTGAGGT 180
Db 123 TGGCCTGTGGCCCGAGGAGCAATGCGTGCCTCAACACCCGGTGCAAGCTTGAGGT 182

Qy 181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGAGCCAG 240
Db 183 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGAGCCAG 242

Qy 241 CCTTGCAGATAAACACACAGAGCTCGGCTCATCGGGAGAAACTGTTCGAGAGTCAG 300
Db 243 CCTTGCAGATAAACACACAGAGCTCGGCTCATCGGGAGAAACTGTTCGAGAGTCAG 302

Qy 301 TGCTAAGGATCAGTGTACCTGTATGAGCAGGCTCTCAACTTCACCTGGAGACATCTCT 360
Db 303 TGCTAAGGATCAGTGTACCTGTATGAGCAGGCTCTCAACTTCACCTGGAGACATCTCT 362

Qy 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTCATATGAGGAGGTGGTGGCTTTCTTGACCAA 420
Db 363 GCTCCCCCAGTCAGACAGGTTCCAGCCCTCATATGAGGAGGTGGTGGCTTTCTTGACCAA 422

Qy 421 ACTGAGCAATCAGCTCAGCTCCTGTGCATCAGTGTGTGACGACGAGCAATCCAGAGAA 480
Db 423 ACTGAGCAATCAGCTCAGCTCCTGTGCATCAGTGTGTGACGACGAGCAATCCAGAGAA 482

Qy 481 TGTGAGAGGCTGAGGAGACAGTGAAGAAAGCTTTGGAGAGCGGAGAGATCAAGCGAT 540
Db 483 TGTGAGAGGCTGAGGAGACAGTGAAGAAAGCTTTGGAGAGCGGAGAGATCAAGCGAT 542

Qy 541 CGGGGAACTGGACCTGCTGTTTATGCTCTGAGAAATGCTTCTGAGCGAGGAGGAGAA 600
Db 543 TGGGGAATGAGCTGCTGTTTATGCTCTGAGAAATGCTTCTGAGCGAGGAGGAG 602

Qy 601 CTAGAAACGAGAACTGCTCTCTTCCCTTCTAAGAAACAAATAGATCCCTGATG 660
Db 603 CTAGAAACGAGAACTGCTCTCTTCCCTTCTAAGAAACAAATAGATCCCTGATG 662

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Qy 661 GACTTTTTTACTAAAGGAAAGTGAAGCTTAAGCTCCACCATCATTTAGAAAGTTTCAT 720
Db 663 GACTTTTTTACTAAAGGAAAGTGAAGCTTAAGCTCCACCATCATTTAGAAAGTTTCAT 722

Qy 721 GAAACCTGGCTCAGTTGAAAGAAATAGTGTCAAGTTGTCATGAGACCCAGAGGTAGA 780
Db 723 GAAACCTGGCTCAGTTGAAAGAAATAGTGTCAAGTTGTCATGAGACCCAGAGGTAGA 782

Qy 781 CTTGATAACCAACAGAGTTTCATTTGACAAATATTTTATGTCATTTGATGATCAACAGAAA 840
Db 783 CTTGATAACCAACAGAGTTTCATTTGACAAATATTTTATGTCATTTGATGATCAACAGAAA 842

Qy 841 AGTATGTACTTTTAAAAAATTTGTTGAAAGGAGGTAACTCTCTCATTTCTTAGAAGAAAAG 900
Db 843 ATATGTACTTTTAAAAAATTTGTTGAAAGGAGGTAACTCTCTCATTTCTTAGAAGAAAAG 902

Qy 901 CCTATGTAACCTTCATTTCCATAACCAATACCTTTATATATATGTAAGTTTATTTATTAAGT 960
Db 903 CCTATGTAACCTTCATTTCCATAACCAATATTTTATATATGTAAGTTTATTTATTAAGT 962

Qy 961 ATACATTTTATTTATGTGTCAGTTTATTAATATGATTTATTTATAGAAAAATTTCTCATG 1020
Db 963 ATACATTTTATTTATGTGTCAGTTTATTAATATGATTTATTTATAGAAAAATTTCTCCTA 1022

Qy 1021 TTGATATTTGAGTATAAGCAAAATATTTATGATAATAAATCACTATAGAAAAACAAGATATCT 1080
Db 1023 TTGATATTT-AGTATAAGGCAAAATATTTATGATAATAAATCACTATGAAAAACAAGATATCT 1081

Qy 1081 TAGCCTTTAATAAACACATCAATATCAAAA 1111
Db 1082 TAGCCTTTAATAAACACATCAATATCAAAA 1112

RESULT 8
US-09-354-243B-7
; Sequence 7, Application US/09354243B
; Patent No. 6359117
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa
; TITLE OF INVENTION: (TIPS)
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.1
; CURRENT APPLICATION NUMBER: US/09/354,243B
; CURRENT FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 7
; LENGTH: 1119
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-09-354-243B-7

Query Match 94.3%; Score 1047.8; DB 3; Length 1119;
Best Local Similarity 97.0%; Pred. No. 5.1e-258;
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;

Qy 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCGGTGATGGCTGTCT 60
Db 3 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCTGTGATGGCTGTCT 62

Qy 61 GCAGAAATCTATAGTTTTTCCCTTATGGGACCTTTGGCCGCCAGCTGCTCTTCAT 120
Db 63 GCAGAAATCTATAGTTTTTCCCTTATGGGACCTTTGGCCGCCAGCTGCTCTTCAT 122

Qy 121 TGGCCTGTGGCCCGAGGAGCAATGCGTGCCTCAACACCCGGTGCAAGCTTGAGGT 180
Db 123 TGGCCTGTGGCCCGAGGAGCAATGCGTGCCTCAACACCCGGTGCAAGCTTGAGGT 182

Qy 181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGAGCCAG 240
Db 183 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGAGCCAG 242

Qy 241 CCTTGCAGATAAACACACAGAGCTCGGCTCATCGGGAGAAACTGTTCGAGAGTCAG 300
Db 243 CCTTGCAGATAAACACACAGAGCTCGGCTCATCGGGAGAAACTGTTCGAGAGTCAG 302

Qy 301 TGCTAAGGATCAGTGTACCTGTATGAGCAGGCTCTCAACTTCACCTGGAGACATCTCT 360
Db 303 TGCTAAGGATCAGTGTACCTGTATGAGCAGGCTCTCAACTTCACCTGGAGACATCTCT 362

Qy 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTCATATGAGGAGGTGGTGGCTTTCTTGACCAA 420
Db 363 GCTCCCCCAGTCAGACAGGTTCCAGCCCTCATATGAGGAGGTGGTGGCTTTCTTGACCAA 422

Qy 421 ACTGAGCAATCAGCTCAGCTCCTGTGCATCAGTGTGTGACGACGAGCAATCCAGAGAA 480
Db 423 ACTGAGCAATCAGCTCAGCTCCTGTGCATCAGTGTGTGACGACGAGCAATCCAGAGAA 482

Qy 481 TGTGAGAGGCTGAGGAGACAGTGAAGAAAGCTTTGGAGAGCGGAGAGATCAAGCGAT 540
Db 483 TGTGAGAGGCTGAGGAGACAGTGAAGAAAGCTTTGGAGAGCGGAGAGATCAAGCGAT 542

Qy 541 CGGGGAACTGGACCTGCTGTTTATGCTCTGAGAAATGCTTCTGAGCGAGGAGGAGAA 600
Db 543 TGGGGAATGAGCTGCTGTTTATGCTCTGAGAAATGCTTCTGAGCGAGGAGGAG 602

Qy 601 CTAGAAACGAGAACTGCTCTCTTCCCTTCTAAGAAACAAATAGATCCCTGATG 660
Db 603 CTAGAAACGAGAACTGCTCTCTTCCCTTCTAAGAAACAAATAGATCCCTGATG 662

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181 GTCCAACTTCCAGCAGCCGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGAGGCCAG 240  
183 GTCCAACTTCCAGCAGCCGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGAGGCCAG 242  
241 CTTTGCAGATAACAACACAGACGCTCCGGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 300  
243 CTTTGCAGATAACAACACAGACGCTCCGGCTCATCGGGGAGAACTGTTCCGAGAGTCAG 302  
301 TGCTAAGGATCAGTGTACCTGTGATGAAGCAGGTGCTCAACTTCAACCTCGGAAGACATCT 360  
303 TGCTAAGATCAGTGTACCTGTGATGAAGCAGGTGCTCAACTTCAACCTCGGAAGACATCT 362  
361 GCTCCCCCACTGACAGAGTTCGGGCCCTACATGCGAGAGGTGGTCTTCTCTGACCAA 420  
363 GCTCCCCCACTGACAGAGTTCGGGCCCTACATGCGAGAGGTGGTCTTCTCTGACCAA 422  
421 ACTCAGCAATCAGTGTACCTGTGATGAAGCAGGTGCTCAACTTCAACCTCGGAAGACATCT 480  
423 ACTCAGCAATCAGTGTACCTGTGATGAAGCAGGTGCTCAACTTCAACCTCGGAAGACATCT 482  
481 TGTCAAGAGCTCAAGAGACAGTGAAGAAAGCTTGGAGAGCGGAGAGATCAAGCGAT 540  
483 TGTCAAGAGCTCAAGAGACAGTGAAGAAAGCTTGGAGAGCGGAGAGATCAAGCGAT 542  
541 CGGGAACTGGACCTGCTGTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAAAG 600  
543 TGGGAATCGACCTGCTGTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAAAG 602  
601 CTGAGAAACGAGAACTGCTCTTCTGCGCTCTTAAAGAAACAATAAGATCCCTGAATG 660  
603 CTGAGAAACGAGAACTGCTCTTCTGCGCTCTTAAAGAAACAATAAGATCCCTGAATG 662  
661 GACTTTTTTAAAGAGAAAGTGAAGAGCTAAAGTCCACCATCATTTAGAGATTTACAT 720  
663 GACTTTTTTAAAGAGAAAGTGAAGAGCTAAAGTCCACCATCATTTAGAGATTTACAT 722  
721 GAAACCTGGCTCAGTTGAAAGAGAAATAGTGTCAAGTTGCTCCATGAGACGAGGTTAG 780  
723 GAAACCTGGCTCAGTTGAAAGAGAAATAGTGTCAAGTTGCTCCATGAGACGAGGTTAG 782  
781 CTTGATAACCAAGATTCATTTGACAAATTTTATTTGTCATTTGATGATAAAGCAAGAAA 840  
783 CTTGATAACCAAGATTCATTTGACAAATTTTATTTGTCATTTGATGATAAAGCAAGAAA 842  
841 AGTATGTAATTTAAAGAAATTTGTTGAAAGAGGTTACCTCTCATCTCTAGAGAAAG 900  
843 ATAATGTAATTTAAAGAAATTTGTTGAAAGAGGTTACCTCTCATCTTTAGAAAGAAAG 902  
901 CCTATGTAATTTCAATTTCCATAACCAATCTTTATATATATGTAAGTTTATTTATATAAGT 960  
903 CTTATGTAATTTCAATTTCCATAACCAATCTTTATATATATGTAAGTTTATTTATATAAGT 962  
961 ATACATTTTATTTATGTCAGTTTATTAATGAGATTTATTTAGAAAAATTTATCTGATG 1020  
963 ATACATTTTATTTATGTCAGTTTATTAATGAGATTTATTTAGAAAAATTTATCTGCTA 1022  
1021 TTGATATTTGAGTATTAAGCAATATATTTATGATAATTAATCTATAGAACAGATATCT 1080  
1023 TTGATATTT-AGTATAGGCAATATATTTATGACAAATTAATCTATGAAACAAGATATCT 1081  
1081 TAGGCTTTTAAATAACACATGAATATCATAAA 1111  
1082 TAGGCTTTTAAATAACACATGAATATCATAAA 1112

RESULT 9

US-09-178-973B-17  
; Sequence 17, Application US/09178973B  
; Patent No. 6274710  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila

RESULT 10

US-09-419-568F-29  
; Sequence 29, Application US/09419568F  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFS) The Proteins Encoded, and Uses Thereof

; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFS)  
; FILE REFERENCE: LUD 5543  
; CURRENT APPLICATION NUMBER: US/09/178, 973B  
; CURRENT FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 17  
; SEQ ID NO 17  
; LENGTH: 5935  
; TYPE: DNA  
; ORGANISM: Mus musculus  
US-09-178-973B-17

Query Match 54.2%; Score 602.4; DB 3; Length 5935;  
Best Local Similarity 99.8%; Pred. No. 1e-143;  
Matches 603; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 508 AAAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGGAACTCGACCTGCTGTTTATGTC 567  
Db 5221 ATAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGGAACTCGACCTGCTGTTTATGTC 5280  
QY 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAACGAGAACTGCTCTCTTCT 627  
Db 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAACGAGAACTGCTCTCTTCT 5340  
QY 628 GCCTTCTTAAAGAAACAATAAGATCCCTGAATGCGACTTTTTTACTAAAGGAAAGTGAGAA 687  
Db 5341 GCCTTCTTAAAGAAACAATAAGATCCCTGATGCGACTTTTTTACTAAAGGAAAGTGAGAA 5400  
QY 688 GCTAACGCTCCACCATCATTTAGAGATTTTCAATGAAACCTGGCTCAGTTGAAAGAGAAAA 747  
Db 5401 GCTAACGCTCCACCATCATTTAGAGATTTTCAATGAAACCTGGCTCAGTTGAAAGAGAAAA 5460  
QY 748 TAGTGTCAAGTTGCTCCATGAGACCGAGGTAGACTTTGATTAACCAAGATTTCAATTGACA 807  
Db 5461 TAGTGTCAAGTTGCTCCATGAGACCGAGGTAGACTTTGATTAACCAAGATTTCAATTGACA 5520  
QY 808 ATATTTTATTTGTCATTTGATAATGCAACAGAAAAAGTATGTACTTTTAAAAAATTTGTTGAA 867  
Db 5521 ATATTTTATTTGTCATTTGATAATGCAACAGAAAAAGTATGTACTTTTAAAAAATTTGTTGAA 5580  
QY 868 AGGAGTTTACCTCTCATTTCTCTAGAGAAAAAGCCTATGTAACCTCATTTTCCATAACCAA 927  
Db 5581 AGGAGTTTACCTCTCATTTCTCTAGAGAAAAAGCCTATGTAACCTCATTTTCCATAACCAA 5640  
QY 928 TACTTTATATATGTAAGTTTATTTATATAGTATACATTTTATTTATGTCAGTTTATTA 987  
Db 5641 TACTTTATATATGTAAGTTTATTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700  
QY 988 ATATGGAATTTTATATAGAAAAATTTATCTGATGTTGATATTTTTCAGTATATAAGCAATAAT 1047  
Db 5701 ATATGGAATTTTATATAGAAAAATTTATCTGATGTTGATATTTTTCAGTATATAAGCAATAAT 5760  
QY 1048 ATTTATGATAATAACTATAGAAACAGATATCTTAGGCTTTTATAAACACATGAATATCA 1107  
Db 5761 ATTTATGATAATAACTATAGAAACAGATATCTTAGGCTTTTATAAACACATGAATATCA 5820  
QY 1108 TAAA 1111  
Db 5821 TAAA 5824

FILE REFERENCE: LUD 5543.2  
CURRENT APPLICATION NUMBER: US/09/419,568F  
CURRENT FILING DATE: 1999-10-18  
PRIOR APPLICATION NUMBER: US09/354,243  
PRIOR FILING DATE: 1999-07-16  
PRIOR APPLICATION NUMBER: US09/178,973  
PRIOR FILING DATE: 1998-10-26  
NUMBER OF SEQ ID NOS: 29  
SEQ ID NO 29  
LENGTH: 5935  
TYPE: DNA  
ORGANISM: Mus musculus  
FEATURE:  
US-09-419-568F-29

Query Match 54.2%; Score 602.4; DB 3; Length 5935;  
Best Local Similarity 99.8%; Pred. No. 1e-143;  
Matches 603; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy	508	AAAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCTGTTATGTC	567
Db	5221	ATAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCTGTTATGTC	5280
Qy	568	TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGAACTGCTCTTCCT	627
Db	5281	TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGAACTGCTCTTCCT	5340
Qy	628	GCCTTCTAAAAGAACCAATAGATCCCTGGAATGCACTTTTACTTAAAGGAAAGTGAGAA	687
Db	5341	GCCTTCTAAAAGAACCAATAGATCCCTGGAATGCACTTTTACTTAAAGGAAAGTGAGAA	5400
Qy	688	GCTAACGTCACCAATAGAGATTTACATGAAACCTGGCTCAGTTGAAAGAGAAAA	747
Db	5401	GCTAACGTCACCAATAGAGATTTACATGAAACCTGGCTCAGTTGAAAGAGAAAA	5460
Qy	748	TAGTGTCAAGTTGTCCATGAGACGAGGTAGACTTGATTAACCAAAAGATTTCATTGACA	807
Db	5461	TAGTGTCAAGTTGTCCATGAGACGAGGTAGACTTGATTAACCAAAAGATTTCATTGACA	5520
Qy	808	ATATTTTATGTCATTGATTAATGCAACAGAAAAAGTATGTAATTTTAAAAAATGTTTGA	867
Db	5521	ATATTTTATGTCATTGATTAATGCAACAGAAAAAGTATGTAATTTTAAAAAATGTTTGA	5580
Qy	868	AGGAGGTTACCTCTCATTCCTCTAGAGAAAGCCTATGTAACCTTTCATTAACCA	927
Db	5581	AGGAGGTTACCTCTCATTCCTCTAGAGAAAGCCTATGTAACCTTTCATTAACCA	5640
Qy	928	TACTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA	987
Db	5641	TACTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA	5700
Qy	988	ATATGGATTTATTTATAGAAAAATTTATCTGATGTTGATATTGAGTATTAAGCAAAATAT	1047
Db	5701	ATATGGATTTATTTATAGAAAAATTTATCTGATGTTGATATTGAGTATTAAGCAAAATAT	5760
Qy	1048	ATTTATGATATAACTATAGAAAAAAGATATCTTAGGCTTTTAAATTAACACATGATATCA	1107
Db	5761	ATTTATGATATAACTATAGAAAAAAGATATCTTAGGCTTTTAAATTAACACATGATATCA	5820
Qy	1108	TAAA 1111	
Db	5821	TAAA 5824	

RESULT 11  
US-09-354-243B-29  
; Sequence 29, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa

TITLE OF INVENTION: (TIFS)  
TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof  
FILE REFERENCE: LUD 5543.1  
CURRENT APPLICATION NUMBER: US/09/354,243B  
CURRENT FILING DATE: 1999-07-16  
PRIOR APPLICATION NUMBER: US09/178,973  
PRIOR FILING DATE: 1998-10-26  
NUMBER OF SEQ ID NOS: 29  
SEQ ID NO 29  
LENGTH: 5935  
TYPE: DNA  
ORGANISM: Homo sapiens  
FEATURE:  
US-09-354-243B-29

Query Match 54.2%; Score 602.4; DB 3; Length 5935;  
Best Local Similarity 99.8%; Pred. No. 1e-143;  
Matches 603; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy	508	AAAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCTGTTATGTC	567
Db	5221	ATAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCTGTTATGTC	5280
Qy	568	TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGAACTGCTCTTCCT	627
Db	5281	TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGAACTGCTCTTCCT	5340
Qy	628	GCCTTCTAAAAGAACCAATAGATCCCTGGAATGCACTTTTACTTAAAGGAAAGTGAGAA	687
Db	5341	GCCTTCTAAAAGAACCAATAGATCCCTGGAATGCACTTTTACTTAAAGGAAAGTGAGAA	5400
Qy	688	GCTAACGTCACCAATAGAGATTTACATGAAACCTGGCTCAGTTGAAAGAGAAAA	747
Db	5401	GCTAACGTCACCAATAGAGATTTACATGAAACCTGGCTCAGTTGAAAGAGAAAA	5460
Qy	748	TAGTGTCAAGTTGTCCATGAGACGAGGTAGACTTGATTAACCAAAAGATTTCATTGACA	807
Db	5461	TAGTGTCAAGTTGTCCATGAGACGAGGTAGACTTGATTAACCAAAAGATTTCATTGACA	5520
Qy	808	ATATTTTATGTCATTGATTAATGCAACAGAAAAAGTATGTAATTTTAAAAAATGTTTGA	867
Db	5521	ATATTTTATGTCATTGATTAATGCAACAGAAAAAGTATGTAATTTTAAAAAATGTTTGA	5580
Qy	868	AGGAGGTTACCTCTCATTCCTCTAGAGAAAGCCTATGTAACCTTTCATTAACCA	927
Db	5581	AGGAGGTTACCTCTCATTCCTCTAGAGAAAGCCTATGTAACCTTTCATTAACCA	5640
Qy	928	TACTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA	987
Db	5641	TACTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA	5700
Qy	988	ATATGGATTTATTTATAGAAAAATTTATCTGATGTTGATATTGAGTATTAAGCAAAATAT	1047
Db	5701	ATATGGATTTATTTATAGAAAAATTTATCTGATGTTGATATTGAGTATTAAGCAAAATAT	5760
Qy	1048	ATTTATGATATAACTATAGAAAAAAGATATCTTAGGCTTTTAAATTAACACATGATATCA	1107
Db	5761	ATTTATGATATAACTATAGAAAAAAGATATCTTAGGCTTTTAAATTAACACATGATATCA	5820
Qy	1108	TAAA 1111	
Db	5821	TAAA 5824	

RESULT 12  
US-09-178-973B-8  
; Sequence 8, Application US/09178973B  
; Patent No. 6274710  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; TITLE OF INVENTION: (TIPS)  
; FILE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543  
; CURRENT APPLICATION NUMBER: US/09/178, 973B  
; CURRENT FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 17  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
US-09-178-973B-8

Query Match 50.0%; Score 555.2; DB 3; Length 7445;  
Best Local Similarity 96.0%; Pred. No. 1.3e-131;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
  
QY 508 AAAGCTTGGAGAGAGCGGAGAGATCAAGCGATCGGGGAAGTGGAGTCTGCTGTTTATGTC 567  
DB 5535 ATAGCTTGGAGAGAGTGGAGAGATCAAGCGATCGGGGAAGTGGAGTCTGCTGTTTATGTC 6594  
  
QY 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAGAACTGCTCCTTCCT 627  
DB 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAGAACTGCTCCTTCCT 6654  
  
QY 628 GCCTTCTAAAAGAAACAATAAGATCCCTGAATGGACTTTTACTAAAGAAAGTGAGAA 687  
DB 6655 GCCTTCTAAAAGAAACAATAAGATCCCTGAATGGACTTTTACTAAAGAAAGTGAGAA 6714  
  
QY 688 GCTAACGCTCCACCATCATTAAGAGATTTCACATGAACCTGGCTCAGTTGAAAGAGAA 747  
DB 6715 GCTAACGCTCCATCATTAAGAGATTTCACATGAACCTGGCTCAGTTGAAAGAGAA 6774  
  
QY 748 TAGTGTCAAGTTGCTCCATGAGACGAGAGGTAGACTTTGATAACCAAGAAAGTTCATTGACA 807  
DB 6775 TAGTGTCAAGTTGCTCCATGAGACGAGAGGTAGACTTTGATAACCAAGAAAGTTCATTGACA 6834  
  
QY 808 ATATTTTATGTCATTGATAATGCAACAGAAAAAGTATGACTTTTAAAAAATGTTTGAA 867  
DB 6835 ATATTTTATGTCATTGATAATGCAACAGAAAAAATAATGACTTTTAAAAAATGTTTGAA 6894  
  
QY 868 AGAGGTTACCTCTCATCTCTAGAGAAAGCCTATGAACTTCATTTCCATACCAA 927  
DB 6895 AGAGGTTACCTCTCATCTCTTAGAAAAAAGCCTATGAACTTCATTTCCATACCAA 6954  
  
QY 928 TACTTTATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 987  
DB 6955 TATTTTATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 7014  
  
QY 988 ATATGGAATTTATATAGAAAAATTAATCTGATGTTGATATTTAGTATATAAGCAAAATAT 1047  
DB 7015 ATATGGAATTTATATAGAAAAATTAATCTGATGTTGATATTT-AGTATAAGGCAAAATAT 7073  
  
QY 1048 ATTTATGATAAATCACTATAGAAACAAGATATCTTAGCTTTAATAACACATGAATATCA 1107  
DB 7074 ATTTATGATAAATCACTATAGAAACAAGATATCTTAGCTTTAATAACACATGAATATCA 7133  
  
QY 1108 TAAA 1111  
DB 7134 TAAA 7137

RESULT 13  
US-09-419-568F-8  
; Sequence 8, Application US/09419568F  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/419, 568F

; CURRENT FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-419-568F-8

Query Match 50.0%; Score 555.2; DB 3; Length 7445;  
Best Local Similarity 96.0%; Pred. No. 1.3e-131;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
  
QY 508 AAAGCTTGGAGAGAGCGGAGAGATCAAGCGATCGGGGAAGTGGAGTCTGCTGTTTATGTC 567  
DB 6535 ATAGCTTGGAGAGAGTGGAGAGATCAAGCGATCGGGGAAGTGGAGTCTGCTGTTTATGTC 6594  
  
QY 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAGAACTGCTCCTTCCT 627  
DB 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAGAACTGCTCCTTCCT 6654  
  
QY 628 GCCTTCTAAAAGAAACAATAAGATCCCTGAATGGACTTTTACTAAAGAAAGTGAGAA 687  
DB 6655 GCCTTCTAAAAGAAACAATAAGATCCCTGAATGGACTTTTACTAAAGAAAGTGAGAA 6714  
  
QY 688 GCTAACGCTCCACCATCATTAAGAGATTTCACATGAACCTGGCTCAGTTGAAAGAGAA 747  
DB 6715 GCTAACGCTCCATCATTAAGAGATTTCACATGAACCTGGCTCAGTTGAAAGAGAA 6774  
  
QY 748 TAGTGTCAAGTTGCTCCATGAGACGAGAGGTAGACTTTGATAACCAAGAAAGTTCATTGACA 807  
DB 6775 TAGTGTCAAGTTGCTCCATGAGACGAGAGGTAGACTTTGATAACCAAGAAAGTTCATTGACA 6834  
  
QY 808 ATATTTTATGTCATTGATAATGCAACAGAAAAAGTATGACTTTTAAAAAATGTTTGAA 867  
DB 6835 ATATTTTATGTCATTGATAATGCAACAGAAAAAATAATGACTTTTAAAAAATGTTTGAA 6894  
  
QY 868 AGAGGTTACCTCTCATCTCTAGAGAAAGCCTATGAACTTCATTTCCATACCAA 927  
DB 6895 AGAGGTTACCTCTCATCTCTTAGAAAAAAGCCTATGAACTTCATTTCCATACCAA 6954  
  
QY 928 TACTTTATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 987  
DB 6955 TATTTTATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 7014  
  
QY 988 ATATGGAATTTATATAGAAAAATTAATCTGATGTTGATATTTAGTATATAAGCAAAATAT 1047  
DB 7015 ATATGGAATTTATATAGAAAAATTAATCTGATGTTGATATTT-AGTATAAGGCAAAATAT 7073  
  
QY 1048 ATTTATGATAAATCACTATAGAAACAAGATATCTTAGCTTTAATAACACATGAATATCA 1107  
DB 7074 ATTTATGATAAATCACTATAGAAACAAGATATCTTAGCTTTAATAACACATGAATATCA 7133  
  
QY 1108 TAAA 1111  
DB 7134 TAAA 7137

RESULT 14  
US-09-354-243B-8  
; Sequence 8, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: The Proteins Encoded, and Uses Thereof

FILE REFERENCE: LUD 5543.1  
CURRENT APPLICATION NUMBER: US/09/354,243B  
CURRENT FILING DATE: 1999-07-16  
PRIOR APPLICATION NUMBER: US09/178,973  
PRIOR FILING DATE: 1998-10-26  
NUMBER OF SEQ ID NOS: 29  
SEQ ID NO 8  
LENGTH: 7445  
TYPE: DNA  
ORGANISM: Mus musculus  
FEATURE:  
US-09-354-243B-8

Query Match 50.0%; Score 555.2; DB 3; Length 7445;  
Best Local Similarity 96.0%; Pred. No. 1.3e-131;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
QY 508 AAGCTTGGAGAGCGGAGATCAAGCGATCGGGAACTGGACCTGCTGTTTATGTC 567  
DB 6535 ATAGCTTGGAGAGTGGAGATCAAGCGATGGGAACTGGACCTGCTGTTTATGTC 6594  
QY 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAAGAACTGCTTTCCT 627  
DB 6595 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAAGAACTGCTTTCCT 6654  
QY 628 GCCTTCTAAAAGAACAAATAGATCCCTGAATGACATTTTAACTAAAGGAAAGTGAGAA 687  
DB 6655 GCCTTCTAAAAGAACAAATAGATCCCTGAATGACATTTTAACTAAAGGAAAGTGAGAA 6714  
QY 688 GCTAACGTCACCATCATTTAGAAATTTCAATGAAACCTGGCTCAGTTGAAAAGAGAAA 747  
DB 6715 GCTAACGTCACCATCATTTAGAAATTTCAATGAAACCTGGCTCAGTTGAAAAGAGAAA 6774  
QY 748 TAGTGTCAAGTTGTCCATGAGACAGAGGTAGACTTGATAAACCAAGAAATTCATTGACA 807  
DB 6775 TAGTGTCAAGTTGTCCATGAGACAGAGGTAGACTTGATAAACCAAGAAATTCATTGACA 6834  
QY 808 ATATTTTATGTCATTCATGATGCAACAGAAAAGTATGACTTTAAAGAAATGTTTGA 867  
DB 6835 ATATTTTATGTCATTCATGATGCAACAGAAAAGTATGACTTTAAAGAAATGTTTGA 6894  
QY 868 AGGAGGTACCTCTCATTCTCTAGAGAAAGCTATGTAACCTTCAATTTCCATAACCAA 927  
DB 6895 AGGAGGTACCTCTCATTCTCTAGAGAAAGCTATGTAACCTTCAATTTCCATAACCAA 6954  
QY 928 TACTTTATATGTAAGTTTATTTTATTAAGTATACATTTTATTAATGTCAGTTTATTA 987  
DB 6955 TATTTTATATGTAAGTTTATTTTATTAAGTATACATTTTATTAATGTCAGTTTATTA 7014  
QY 988 ATATGGATTTATTTATAGAAAATTTCTGATGTTGATATTTGAGTATTAAGCAATAAT 1047  
DB 7015 ATATGGATTTATTTATAGAAAATTTCTGATGTTGATATTTTATTAAGCAATAAT 7073  
QY 1048 ATTTATGATAATACTATAGAAAACAGATATCTTAGGCTTTAAATAACACATGATATCA 1107  
DB 7074 ATTTATGATAATACTATAGAAAACAGATATCTTAGGCTTTATTAAGCAATAATGATATCA 7133  
QY 1108 TAAA 1111  
DB 7134 TAAA 7137

RESULT 15  
US-10-084-298-1  
Sequence 1, Application US/10084298  
Patent No. 6939545  
GENERAL INFORMATION:  
APPLICANT: Jacobs, Kenneth  
APPLICANT: Pittman, Debra  
APPLICANT: Fouser, Lynette  
APPLICANT: Spaulding, Vikki  
APPLICANT: Xuan, Dejun  
TITLE OF INVENTION: Composition and Method for Treating Inflammatory

TITLE OF INVENTION: Disorders  
FILE REFERENCE: G15358 CIP  
CURRENT APPLICATION NUMBER: US/10/084,298  
CURRENT FILING DATE: 2002-09-10  
PRIOR APPLICATION NUMBER: 60/270,823  
PRIOR FILING DATE: 2001-02-23  
PRIOR APPLICATION NUMBER: 60/281,353  
PRIOR FILING DATE: 2001-04-03  
PRIOR APPLICATION NUMBER: 60/131,473  
PRIOR FILING DATE: 1999-04-28  
PRIOR APPLICATION NUMBER: 09/561,811  
PRIOR FILING DATE: 2000-04-28  
NUMBER OF SEQ ID NOS: 10  
SOFTWARE: PatentIn Ver. 2.1  
SEQ ID NO 1  
LENGTH: 1191  
TYPE: DNA  
ORGANISM: Homo sapiens  
US-10-084-298-1  
Query Match 47.6%; Score 529; DB 3; Length 1191;  
Best Local Similarity 72.7%; Pred. No. 2.9e-125;  
Matches 830; Conservative 0; Mismatches 275; Indels 36; Gaps 10;  
QY 7 CTCTCCTCTCAGTTATCAACTTTTGCACCTTGTGCGATCGGTGATGGCTGTCTCTGCAGAA 66  
DB 28 CTCTTCCCAGTCACCACTTGTCTCGAGTTAGAAATTTCTGCAATGGCCGCTGCAGAA 87  
QY 67 ATCTATGAGTTTTCCTTTATGGGACTTTTGGCGGCGAGCTGCTGCTTCTCATTTGCCCT 126  
DB 88 ATCTGAGCTCTTCTTATGGGACCTTGGCCGCTGCACAGCTGCTCTTCTTGGCCCT 147  
QY 127 GTGGGCCAGAGCAAAATGGCTGCCCATCAACCCGCTGCAAGCTTCAGGTGTCCAA 186  
DB 148 CTGGTACAGGAGGAGCAGCTGCGCCCATCAGTCCCACTGAGGCTTGCAAGTCCAA 207  
QY 187 CTTCACAGCCGCTACATCTCAACCGCACCTTTATGCTGCCCAAGAGGCGCAGCTTGC 246  
DB 208 CTTCACAGCCCTATATACCAACCGCACCTTCATGCTGCTTAAGGAGCTAGTTGGC 267  
QY 247 AGATAACAAACAGAGCTCGGCTCATCGGGGAGAACTGTTCCGAGAGTCAAGTCTAA 306  
DB 268 TGATAACAAACAGAGCTGCTCTCATTTGGGGAGAACTGTTCCAGGAGTCAATGAG 327  
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QY 367 CCAGTCAGACAGGTTCCGGCCCTACATGACGAGGCTGCTGCTTCTTCTGACCAACTCAG 426  
DB 388 TCAATCTGATAGTTTCCAGCTTATATGAGGAGGTTGCTGCTTCTTCTGCGCAGCTCAG 447  
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QY 487 AAGCTGAAGAGACAGTGAAGAAAGCTTGAAGAGCGGAGAGATCAAGCGATCGGGA 546  
DB 508 AAAGCTGAAGGACACAGTGAAGAAAGCTTGAAGAGGAGTGAAGCAATTTGGAGA 567  
QY 547 ACTGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCAGAGAAAGCTAGAA 606  
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DB 628 AATGAATAACTAACCCCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 687  
QY 667 TTTA-----CTAAGGAAAGTGAAGTCAAGTCAAGTCAAGTCAAGTCAAGTCAAGTCA 722  
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QY 723 AACCTGCTCAGTTGAAAGAGAAATAGTGTCAA--GTTGTCCATGAGACCAG-AGGTAG 779



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GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

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(without alignments)  
16534.837 Million cell updates/sec

Title: US-09-751-797-9

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Gapop 10.0 , Gapext 1.0

Searched: 9793542 seqs, 4134689005 residues

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Maximum DB seq length: 2000000000

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Maximum Match 100%

Listing first 45 summaries

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- 10: /cgn2\_6/ptodata/1/pubpna/US11\_PUBCOMB.seq:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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2	1111	100.0	1111	7	US-10-627-273-9
3	1049.4	94.5	1166	5	US-10-084-298-3
4	1049.4	94.5	1166	6	US-10-256-977-3
5	1049.4	94.5	1166	8	US-10-873-972-3
6	1049.4	94.5	1166	10	US-11-157-387-3
7	1048.4	94.4	1050	5	US-10-090-365-40
8	1048.4	94.4	1050	5	US-10-104-919-42
9	1048.4	94.4	1050	8	US-10-807-837-10
10	1048.4	94.4	1050	9	US-10-968-432-42
11	1048.4	94.4	1050	10	US-11-045-944-40
12	1047.8	94.3	1119	3	US-09-751-797-7
13	1047.8	94.3	1119	7	US-10-627-273-7
14	757.2	68.2	778	3	US-09-746-375-37
15	757.2	68.2	778	7	US-10-395-741B-37
16	757.2	68.2	778	7	US-10-806-294-37
17	602.4	54.2	5935	3	US-09-751-797-29
18	602.4	54.2	5935	7	US-10-627-273-29
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20	555.2	50.0	7445	7	US-10-627-273-8
21	529	47.6	1177	10	US-11-013-741-1
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25	529	47.6	1191	8	US-10-873-972-1	Sequence 1, Appli
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27	525.6	47.3	1152	3	US-09-870-574-1	Sequence 1, Appli
28	525.6	47.3	1152	5	US-10-006-867-153	Sequence 153, App
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38	525.6	47.3	1152	5	US-10-063-549-153	Sequence 153, App
39	525.6	47.3	1152	5	US-10-227-884-243	Sequence 243, App
40	525.6	47.3	1152	5	US-10-002-796-125	Sequence 125, App
41	525.6	47.3	1152	5	US-10-066-273-125	Sequence 125, App
42	525.6	47.3	1152	5	US-10-066-494-125	Sequence 125, App
43	525.6	47.3	1152	5	US-10-230-163-243	Sequence 243, App
44	525.6	47.3	1152	5	US-10-063-554-153	Sequence 153, App
45	525.6	47.3	1152	5	US-10-066-269-125	Sequence 125, App

#### ALIGNMENTS

##### RESULT 1

US-09-751-797-9

; Sequence 9, Application US/09751797

; Patent No. US20010024652A1

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Leuhed, Jamila

; APPLICANT: Renaud, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; FILE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5543.2

; CURRENT APPLICATION NUMBER: US/09/751,797

; CURRENT FILING DATE: 2000-12-29

; PRIOR APPLICATION NUMBER: 09/419,568

; PRIOR FILING DATE: 1999-10-18

; PRIOR APPLICATION NUMBER: US09/178,973

; PRIOR FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 29

; SEQ ID NO 9

; LENGTH: 1111

; TYPE: DNA

; ORGANISM: Mus musculus

; FEATURE:

US-09-751-797-9

Query Match 100.0%; Score 1111; DB 3; Length 1111;  
Best Local Similarity 100.0%; Pred. No. 7.4e-229;  
Matches 1111; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY	1	AACAGGCTCTCTCTCAGTTATCACTTTTGACACTTGTGCGATCGGTGATGGCTGTCCT	60
Db	1	AACAGGCTCTCTCTCAGTTATCACTTTTGACACTTGTGCGATCGGTGATGGCTGTCCT	60
QY	61	GCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTGGCCGCGAGTGCCTGCTTCAT	120
Db	61	GCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTGGCCGCGAGTGCCTGCTTCAT	120
QY	121	TGCCCTGTGGCCCGCAGGACAAATGCGTGCCTATCAACACCGGTGCAAGCTTGAGGT	180
Db	121	TGCCCTGTGGCCCGCAGGACAAATGCGTGCCTATCAACACCGGTGCAAGCTTGAGGT	180
QY	181	GTCCTAACTTCCACGACCGGTACATGTCACCCACCTTTATGCTGSCCAAGGAGCCAG	240
Db	181	GTCCTAACTTCCACGACCGGTACATGTCACCCACCTTTATGCTGSCCAAGGAGCCAG	240
QY	241	CTTGCAGATAACAACACAGACGTCGGGCTCATCGGGGAGAAACTGTTCCGAGGAGTCAG	300



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Qy 541 CGGGGAACAGTGCCTCCTTCTGAGAAATGCTTCTGAGAAATGCTTCTGAGGAGAGAAAG 600
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Qy 721 GAAACCTGGCTCAGTGTGAAGCAGAAATAGTGTCAAGTTGTCCATGACGACGAGGTAGA 780
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Qy 781 CTTGATAACCAAGATTTCAATGACAATATTTTATTTGTCATTTGATTAATGCAACAGAAAA 840
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Qy 961 ATACATTTTATTTATGTAGTATTAATATGGAATTTATTTATAGAAAAATTTATCTGATG 1020
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Qy 1021 TTGATATTTGATATAAGCAAAATATATTTATGATATAATATATATAGAAACAGATATCT 1080
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; Sequence 9, Application US/10627273
; Publication No. US20040110189A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; FILE OF INVENTION: (ifIs) The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.2
; CURRENT APPLICATION NUMBER: US/10/627,273
; CURRENT FILING DATE: 2003-07-25
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; PRIOR APPLICATION NUMBER: US/09/751,797
; PRIOR FILING DATE: 2000-12-29
; PRIOR APPLICATION NUMBER: 09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 9
; LENGTH: 1111
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-10-627-273-9

Query Match 100.0%; Score 1111; DB 7; Length 1111;
Best Local Similarity 100.0%; Pred. No. 7.4e-229;
Matches 1111; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AACAGGCTCTCCTCTCAGTTTATCAACTTTTGGACACTTTGTGCGATCGGTGATGGCTGTCTCT 60
Db 1 AACAGGCTCTCCTCTCAGTTTATCAACTTTTGGACACTTTGTGCGATCGGTGATGGCTGTCTCT 60
Qy 61 GCAGAAATCTATGAGTTTTTCCCTTATGCGGACTTTTGGCCGCGAGCTGCCCTGTCTTCAT 120
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Qy 121 TGGCCTGTGGCCGAGGAGCAATGCGTCCCATCAACACCGGTGCAAGCTTGAGGT 180
Db 121 TGGCCTGTGGCCGAGGAGCAATGCGTCCCATCAACACCGGTGCAAGCTTGAGGT 180
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Qy 361 GCTCCCCCAGTCACAGAGTTCCGGCCCTACATGACGAGGTGCTGCTTTCCTGACCAA 420
Db 361 GCTCCCCCAGTCACAGAGTTCCGGCCCTACATGACGAGGTGCTGCTTTCCTGACCAA 420
Qy 421 ACTCAGCAATCAGCTCAGCTCCTGTCTCATCAGTGGTGACGACCAAGACATCCAGAGAA 480
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Qy 481 TGTCAAGGCTGAAGCAGTGAAGAAAGCTTGGAGAGGCGGAGAGATCAAAAGCAT 540
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; FILE REFERENCE: GI5358 CIP
; CURRENT APPLICATION NUMBER: US/10/256,977
; CURRENT FILING DATE: 2002-09-27
; PRIOR APPLICATION NUMBER: US/10/084,298
; PRIOR FILING DATE: 2002-09-10
; PRIOR APPLICATION NUMBER: 60/270,823
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/281,353
; PRIOR FILING DATE: 2001-04-03
; PRIOR APPLICATION NUMBER: 60/131,473
; PRIOR FILING DATE: 1999-04-28
; PRIOR APPLICATION NUMBER: 09/561,811
; PRIOR FILING DATE: 2000-04-28
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn ver. 2.1
; SEQ ID NO 3
; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Murine
US-10-256-977-3

Query Match      94.5%; Score 1049.4; DB 6; Length 1166;
Best Local Similarity 97.1%; Pred. No. 1.4e-215;
Matches 1079; Conservative 0; Mismatches 31; Indels 1; Gaps 1;

Qy 1 AACAGGCTCTCCTCAGTATCAACTTTTGACACTTGTGCGATCGGTGATGGCTGCT 60
Db 26 AACAGGCTCTCCTCAGTATCAACTTTTGACACTTGTGCGATCTCTGATGGCTGCT 85

Qy 61 GCAGAAATCTATGATGTTTTTCCCTTATGGGACATTTGGCCGCCAGCTGCCCTTCAT 120
Db 86 GCAGAAATCTATGATGTTTTTCCCTTATGGGACATTTGGCCGCCAGCTGCCCTTCAT 145

Qy 121 TGCCTCTGGGCCAGGAGCAATGCGCTGCCATCAACACCCGGTGCAGCTTGAGGT 180
Db 146 TGCCTCTGGGCCAGGAGCAATGCGCTGCCATCAACACCCGGTGCAGCTTGAGGT 205

Qy 181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240
Db 206 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 265

Qy 241 CTTTGCAGATAAACACACAGACGTCCTGCTCATCGGGAGAGAACTGTTCCGAGGAGTCAG 300
Db 266 CTTTGCAGATAAACACACAGATGTCGGCTCATCGGGAGAGAACTGTTCCGAGGAGTCAG 325

Qy 301 TGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACCTCGGAAGACATCT 360
Db 326 TGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACCTCGGAAGACATCT 385

Qy 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCAGGAGGTGGTGGCTTTCTTGACCAA 420
Db 386 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCAGGAGGTGGTGGCTTTCTTGACCAA 445

Qy 421 ACTCAGCAATCAGTCTCAGTCTCTGTCAATCAGTGGTGACGACAGCAACATCCAGAGAA 480
Db 446 ACTCAGCAATCAGTCTCAGTCTCTGTCAATCAGCGGTGACGCCAGCAACATCCAGAGAA 505

Qy 481 TGTCAAGAGGTGAAGAGAGACAGTGAAAGAAAGCTTGGAGAGAGCGGAGAGATCAAGCGAT 540
Db 506 TGTCAAGAGGTGAAGAGAGACAGTGAAAGAAAGCTTGGAGAGAGTGGAGAGATCAAGCGAT 565

Qy 541 CGGGAACTGGACCTGCTGTTTATGCTCTCAGAAATGCTTGGCTCTGAGCGAGAGAGAG 600
Db 566 TGGGAACTGGACCTGCTGTTTATGCTCTCAGAAATGCTTGGCTCTGAGCGAGAGAGAG 625

Qy 601 CTAGAAAACGAGAACTGCTCTCTTCCCTTCTAAAAAGAAACAATAAGATCCCTGATG 660
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Qy 781 CTTGATAAACCAAGAGATTCATTGACAAATATTTTATTTGTCATTTGATGATCAACAGAAA 840
Db 806 CTTGATAAACCAAGAGATTCATTGACAAATATTTTATTTGTCATTTGATGATCAACAGAAA 865

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Qy 1021 TTGATATTTTGCAGTAAAGCAAAATAATATTTATGATAATAAATATAGAAAAACAAGATATCT 1080
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Qy 1081 TAGGCTTTTATAAACACATCATATATATAAA 1111
Db 1105 TAGGCTTTTATAAACACATCATATATATAAA 1135

RESULT 5
US-10-873-972-3
; Sequence 3, Application US/10873972
; Publication No. US20050042220A1
; GENERAL INFORMATION:
; APPLICANT: Li, Jing
; APPLICANT: Tan, Xiang-Yang
; APPLICANT: Tomkinson, Kathleen N.
; APPLICANT: Pittman, Debra D.
; APPLICANT: Veldman, Geertruida M.
; APPLICANT: Fouser, Lynette
; TITLE OF INVENTION: Antibodies Against Interleukin-22 and Uses Therefor
; FILE REFERENCE: AM101524
; CURRENT APPLICATION NUMBER: US/10/873,972
; CURRENT FILING DATE: 2004-06-22
; PRIOR APPLICATION NUMBER: US 60/480,652
; PRIOR FILING DATE: 2003-06-23
; PRIOR APPLICATION NUMBER: US 10/084,298
; PRIOR FILING DATE: 2002-02-25
; PRIOR APPLICATION NUMBER: US 60/270,823
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: US 60/281,353
; PRIOR FILING DATE: 2001-04-03
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Mus musculus
US-10-873-972-3

Query Match      94.5%; Score 1049.4; DB 8; Length 1166;
Best Local Similarity 97.1%; Pred. No. 1.4e-215;
Matches 1079; Conservative 0; Mismatches 31; Indels 1; Gaps 1;

Qy 1 AACAGGCTCTCCTCAGTATCAACTTTTGACACTTGTGCGATCGGTGATGGCTGCT 60
Db 26 AACAGGCTCTCCTCAGTATCAACTTTTGACACTTGTGCGATCTCTGATGGCTGCT 85

Qy 61 GCAGAAATCTATGATGTTTTTCCCTTATGGGACATTTGGCCGCCAGCTGCCCTTCAT 120
Db 86 GCAGAAATCTATGATGTTTTTCCCTTATGGGACATTTGGCCGCCAGCTGCCCTTCAT 145

Qy 121 TGCCTCTGGGCCAGGAGCAATGCGCTGCCATCAACACCCGGTGCAGCTTGAGGT 180
Db 146 TGCCTCTGGGCCAGGAGCAATGCGCTGCCATCAACACCCGGTGCAGCTTGAGGT 205

Qy 181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240
Db 206 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 265

Qy 241 CTTTGCAGATAAACACACAGACGTCCTGCTCATCGGGAGAGAACTGTTCCGAGGAGTCAG 300
Db 266 CTTTGCAGATAAACACACAGATGTCGGCTCATCGGGAGAGAACTGTTCCGAGGAGTCAG 325

Qy 301 TGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACCTCGGAAGACATCT 360
Db 326 TGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACCTCGGAAGACATCT 385

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Qy 481 TGTCAAGAGGTGAAGAGAGACAGTGAAAGAAAGCTTGGAGAGAGCGGAGAGATCAAGCGAT 540
Db 506 TGTCAAGAGGTGAAGAGAGACAGTGAAAGAAAGCTTGGAGAGAGTGGAGAGATCAAGCGAT 565

Qy 541 CGGGAACTGGACCTGCTGTTTATGCTCTCAGAAATGCTTGGCTCTGAGCGAGAGAGAG 600
Db 566 TGGGAACTGGACCTGCTGTTTATGCTCTCAGAAATGCTTGGCTCTGAGCGAGAGAGAG 625

Qy 601 CTAGAAAACGAGAACTGCTCTCTTCCCTTCTAAAAAGAAACAATAAGATCCCTGATG 660
Db 626 CTAGAAAACGAGAACTGCTCTCTTCCCTTCTAAAAAGAAACAATAAGATCCCTGATG 685

Qy 661 GACTTTTTTACTAAAGGAAAGTGAGAGAGCTAACGTCACCATATTAAGAAAGTTTCAT 720
Db 686 GACTTTTTTACTAAAGGAAAGTGAGAGAGCTAACGTCACCATATTAAGAAAGTTTCAT 745
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Db 146 TGGCCCTGTGGCCAGGAGGAAATGCGTCCCGTCAACACCGGTCAGAGCTTGAGGT 205  
Qy 181 GTCCAACTTCAGCAGCGCGTACATCGTCAACCGCACCTTTATGTGGCCCAAGAGGCGAG 240  
Db 206 GTCCAACTTCAGCAGCGCGTACATCGTCAACCGCACCTTTATGTGGCCCAAGAGGCGAG 265  
Qy 241 CTTTGAGATAAACAACAAGAGGTCGGCTCATCGGGGAGAACTGTTCGAGAGGTCAG 300  
Db 266 CTTTGAGATAAACAACAAGAGGTCGGCTCATCGGGGAGAACTGTTCGAGAGGTCAG 325  
Qy 301 TGTCTAAGGATCAGTGTACCTGTAGAGCAGGTCCTCAACTTCACTTCACTGGAAGCATTTCT 360  
Db 326 TGTCTAAGGATCAGTGTACCTGTAGAGCAGGTCCTCAACTTCACTTCACTGGAAGCATTTCT 385  
Qy 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCGAGAGGTGGTGTCTTCTTGACCAA 420  
Db 386 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCGAGAGGTGGTGTCTTCTTGACCAA 445  
Qy 421 ACTCAGCAATCAGCTCAGCTCCTGTGTACATCAGTGTGTGACGACGAGCAATCCAGAGAA 480  
Db 446 ACTCAGCAATCAGCTCAGCTCCTGTGTACATCAGTGTGTGACGACGAGCAATCCAGAGAA 505  
Qy 481 TGTCTAAGGATCAGTGTACCTGTAGAGCAGGTCCTCAACTTCACTTCACTGGAAGCATTTCT 540  
Db 506 TGTCTAAGGATCAGTGTACCTGTAGAGCAGGTCCTCAACTTCACTTCACTGGAAGCATTTCT 565  
Qy 541 CGGGGAAGTGGAGTGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 600  
Db 566 TGGGGAAGTGGAGTGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 625  
Qy 601 CTAGAAAACGAGAACTGCTCTTCTGCTCTTCTGAGAAATGCTTCTGAGAAATGCTTCTGAGAA 660  
Db 626 CTAGAAAACGAGAACTGCTCTTCTGCTCTTCTGAGAAATGCTTCTGAGAAATGCTTCTGAGAA 685  
Qy 661 GACTTTTTTAAAGAAAGTGAAGAGCTTAAAGTGAAGAGCTTAAAGTGAAGAGCTTAAAGTGAAG 720  
Db 686 GACTTTTTTAAAGAAAGTGAAGAGCTTAAAGTGAAGAGCTTAAAGTGAAGAGCTTAAAGTGAAG 745  
Qy 721 GAAACCTGGCTCAGTGTGAAAGAGAAATAGTGTCAAGTGTGTCATGAGACGAGGAGTGA 780  
Db 746 GAAACCTGGCTCAGTGTGAAAGAGAAATAGTGTCAAGTGTGTCATGAGACGAGGAGTGA 805  
Qy 781 CTTGATAACCAAGAGTTCATTGACAAATTTTATTTGTCATTGATTAATGCAACAGAGAA 840  
Db 806 CTTGATAACCAAGAGTTCATTGACAAATTTTATTTGTCATTGATTAATGCAACAGAGAA 865  
Qy 841 AGTATGATCTTTAAAGAAATTTGTTGAAAGAGGTTACCTCTCATTTCTTAGAAGAAAG 900  
Db 866 ATAATGATCTTTAAAGAAATTTGTTGAAAGAGGTTACCTCTCATTTCTTAGAAGAAAG 925  
Qy 901 CCTATGTAATCTTCAATTTCCATACCAATATCTTTATATATGTAAGTTTATTTATTAAGT 960  
Db 926 CTTATGTAATCTTCAATTTCCATACCAATATTTATATATGTAAGTTTATTTATTAAGT 985  
Qy 961 ATACATTTTATTTATGTCAGTTTATTAATATGATTTTATTTATAGAAAATTTATCTGATG 1020  
Db 986 ATACATTTTATTTATGTCAGTTTATTAATATGATTTTATTTATAGAAAATTTATCTGCTA 1045  
Qy 1021 TTGATATTTGAGTATAAGCAATATATTTATGATTAATTAATCTATAGAAAATTTATCTGCTA 1080  
Db 1046 TTGATATTT-AGTATAGGCAATATATTTATGATTAATTAATCTATAGAAAATTTATCTGCTA 1104  
Qy 1081 TAGGCTTTTAAACACATGAATATCATATA 1111  
Db 1105 TAGGCTTTTAAACACATGAATATCATATA 1135

RESULT 6  
US-11-157-387-3  
; Sequence 3, Application US/11157387  
; Publication No. US20050238649A1  
; GENERAL INFORMATION:

; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: Xuan, Dejun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory  
; TITLE OF INVENTION: Disorders  
; FILE REFERENCE: G15358 CIP  
; CURRENT FILING DATE: 2005-06-20  
; PRIOR FILING DATE: 2002-09-10  
; PRIOR FILING DATE: 2002-09-10  
; PRIOR FILING DATE: 2002-09-10  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR FILING DATE: 2001-02-23  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR FILING DATE: 2001-04-03  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR FILING DATE: 1999-04-28  
; PRIOR FILING DATE: 2000-04-28  
; NUMBER OF SEQ ID NOS: 10  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 3  
; LENGTH: 1166  
; TYPE: DNA  
; ORGANISM: Murine  
US-11-157-387-3

Query Match 94.5%; Score 1049.4; DB 10; Length 1166;  
Best Local Similarity 97.1%; Pred. No. 1.4e-215;  
Matches 1079; Conservative 0; Mismatches 31; Indels 1; Gaps 1;  
Qy 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTCGATCGTGTGCTGCT 60  
Db 26 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTCGATCGTGTGCTGCT 85  
Qy 61 GCAGAAATCTATGAGTTTTTCCCTTATGAGGACTTTGGCGCCAGTGTCTTCTCAT 120  
Db 86 GCAGAAATCTATGAGTTTTTCCCTTATGAGGACTTTGGCGCCAGTGTCTTCTCAT 145  
Qy 121 TGGCCTGTGGCCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 180  
Db 146 TGGCCTGTGGCCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 205  
Qy 181 GTCCAACTTCAGCAGCGCGTACATCGTCAACCGCACCTTTATGTGGCCCAAGAGGCGAG 240  
Db 206 GTCCAACTTCAGCAGCGCGTACATCGTCAACCGCACCTTTATGTGGCCCAAGAGGCGAG 265  
Qy 241 CTTTGAGATAAACAACAAGAGGTCGGCTCATCGGGGAGAACTGTTCGAGAGGTCAG 300  
Db 266 CTTTGAGATAAACAACAAGAGGTCGGCTCATCGGGGAGAACTGTTCGAGAGGTCAG 325  
Qy 301 TGTCTAAGGATCAGTGTACCTGTAGAGCAGGTCCTCAACTTCACTTCACTGGAAGCATTTCT 360  
Db 326 TGTCTAAGGATCAGTGTACCTGTAGAGCAGGTCCTCAACTTCACTTCACTGGAAGCATTTCT 385  
Qy 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCGAGAGGTGGTGTCTTCTTGACCAA 420  
Db 386 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCGAGAGGTGGTGTCTTCTTGACCAA 445  
Qy 421 ACTCAGCAATCAGCTCAGCTCCTGTGTACATCAGTGTGTGACGACGAGCAATCCAGAGAA 480  
Db 446 ACTCAGCAATCAGCTCAGCTCCTGTGTACATCAGTGTGTGACGACGAGCAATCCAGAGAA 505  
Qy 481 TGTCTAAGGATCAGTGTACCTGTAGAGCAGGTCCTCAACTTCACTTCACTGGAAGCATTTCT 540  
Db 506 TGTCTAAGGATCAGTGTACCTGTAGAGCAGGTCCTCAACTTCACTTCACTGGAAGCATTTCT 565  
Qy 541 CGGGGAAGTGGAGTGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 600  
Db 566 TGGGGAAGTGGAGTGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 625  
Qy 601 CTAGAAAACGAGAACTGCTCTTCTGCTCTTCTGAGAAATGCTTCTGAGAAATGCTTCTGAGAA 660



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; APPLICANT: Presnell, Scott R.
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Chen, Zhi
; APPLICANT: Hughes, Steven D.
; TITLE OF INVENTION: Human Cytokine Receptor
; FILE REFERENCE: 01-12
; CURRENT APPLICATION NUMBER: US/10/104,919
; CURRENT FILING DATE: 2002-03-23
; PRIOR APPLICATION NUMBER: US 60/279,222
; PRIOR FILING DATE: 2001-03-27
; NUMBER OF SEQ ID NOS: 62
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 42
; LENGTH: 1050
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (5)...(589)
US-10-104-919-42

Query Match          94.4%; Score 1048.4; DB 5; Length 1050;
Best Local Similarity 99.9%; Pred. No. 2.2e-215;
Matches 1049; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCGGTGATGCTGTCT 60
DB 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCGGTGATGCTGTCT 60
QY 61 GCAGAAATCTATGAGTTTTCCCTTATGCGGACTTTTGCGCGCAGCTGCTTCTCAT 120
DB 61 GCAGAAATCTATGAGTTTTCCCTTATGCGGACTTTTGCGCGCAGCTGCTTCTCAT 120
QY 121 TSCCTCTGTGGCCGAGGAGCAATGCGCTGCCATCAACCCGGTGCAAGTTCAGGT 180
DB 121 TSCCTCTGTGGCCGAGGAGCAATGCGCTGCCATCAACCCGGTGCAAGTTCAGGT 180
QY 181 GTCCAACTTCAGGAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240
DB 181 GTCCAACTTCAGGAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240
QY 241 CCTTGAGATAAACAACAGAGCTCGGCTCATCGGGGAGAAACTGTTCGAGAGTTCAG 300
DB 241 CCTTGAGATAAACAACAGAGCTCGGCTCATCGGGGAGAAACTGTTCGAGAGTTCAG 300
QY 301 TGCTAAGGATCAGTGTCTACCTGATGAGAGGCTCACTTCACTTCCCTGGAACATTTCT 360
DB 301 TGCTAAGGATCAGTGTCTACCTGATGAGAGGCTCACTTCACTTCCCTGGAACATTTCT 360
QY 361 GCTCCCCAGTCAAGAGGTTCCGGCCCTACATGACGAGGTGGTGGCTTTCTTGACCAA 420
DB 361 GCTCCCCAGTCAAGAGGTTCCGGCCCTACATGACGAGGTGGTGGCTTTCTTGACCAA 420
QY 421 ACTCAGCAATCAGTCACTGCTCTGTCACTCAGTGTGACGACCCAGAACATCCAGAGAA 480
DB 421 ACTCAGCAATCAGTCACTGCTCTGTCACTCAGTGTGACGACCCAGAACATCCAGAGAA 480
QY 481 TGTGAGAGGCTGAGAGAGAGAGTGTGAGAGAGAGTGTGAGAGAGAGAGTGTGAGAGAG 540
DB 481 TGTGAGAGGCTGAGAGAGAGAGTGTGAGAGAGAGTGTGAGAGAGAGAGTGTGAGAGAG 540
QY 541 CGGGGAGTGGAGTGTGTTTATGCTCTGAGAAATGCTTGGCTGTGAGGAGAGAGAG 600
DB 541 CGGGGAGTGGAGTGTGTTTATGCTCTGAGAAATGCTTGGCTGTGAGGAGAGAGAG 600
QY 601 CTAGAAAACGAGAACTGCTCTTCTGCTTCTTAAAGAAACAATAAGATCCCTGAATG 660
DB 601 CTAGAAAACGAGAACTGCTCTTCTGCTTCTTAAAGAAACAATAAGATCCCTGAATG 660
QY 661 GACTTTTTTCTAAGGAAAGTGAAGAGCTAAAGTCCACCATCATTTAGAAAGATTTCAT 720
DB 661 GACTTTTTTCTAAGGAAAGTGAAGAGCTAAAGTCCACCATCATTTAGAAAGATTTCAT 720

RESULT 9
US-10-807-837-10
; Sequence 10, Application US/10807837
; Publication No. US20040209330A1
; GENERAL INFORMATION:
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Chandrasekhar, Yasmin A.
; APPLICANT: Dillon, Stacey R.
; APPLICANT: Lehner, Joyce M.
; APPLICANT: Siadak, Anthony W.
; APPLICANT: Sivakumar, Pallavur V.
; APPLICANT: Moore, Margaret D.
; TITLE OF INVENTION: ANTI-IL-22RA ANTIBODIES AND BINDING
; PARTNERS AND METHODS OF USING IN INFLAMMATION
; FILE REFERENCE: 03-02
; CURRENT APPLICATION NUMBER: US/10/807,837
; CURRENT FILING DATE: 2004-03-24
; PRIOR APPLICATION NUMBER: US 60/457,481
; PRIOR FILING DATE: 2003-03-24
; PRIOR APPLICATION NUMBER: US 60/523,295
; PRIOR FILING DATE: 2003-11-17
; NUMBER OF SEQ ID NOS: 62
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 1050
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (5)...(589)
US-10-807-837-10

Query Match          94.4%; Score 1048.4; DB 8; Length 1050;
Best Local Similarity 99.9%; Pred. No. 2.2e-215;
Matches 1049; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCGGTGATGCTGTCT 60
DB 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCGGTGATGCTGTCT 60
QY 61 GCAGAAATCTATGAGTTTTCCCTTATGCGGACTTTTGCGCGCAGCTGCTTCTCAT 120
DB 61 GCAGAAATCTATGAGTTTTCCCTTATGCGGACTTTTGCGCGCAGCTGCTTCTCAT 120
QY 121 TGCCCTGTGGCCGAGGAGCAATGCGCTGCCATCAACCCGGTGCAAGTTCAGGT 180
DB 121 TGCCCTGTGGCCGAGGAGCAATGCGCTGCCATCAACCCGGTGCAAGTTCAGGT 180
QY 181 GTCCAACTTCAGGAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240
DB 181 GTCCAACTTCAGGAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240
QY 241 CCTTGAGATAAACAACAGAGCTCGGCTCATCGGGGAGAAACTGTTCGAGAGTTCAG 300
DB 241 CCTTGAGATAAACAACAGAGCTCGGCTCATCGGGGAGAAACTGTTCGAGAGTTCAG 300
QY 301 TGCTAAGGATCAGTGTCTACCTGATGAGAGGCTCACTTCACTTCCCTGGAACATTTCT 360
DB 301 TGCTAAGGATCAGTGTCTACCTGATGAGAGGCTCACTTCACTTCCCTGGAACATTTCT 360
QY 361 GCTCCCCAGTCAAGAGGTTCCGGCCCTACATGACGAGGTGGTGGCTTTCTTGACCAA 420
DB 361 GCTCCCCAGTCAAGAGGTTCCGGCCCTACATGACGAGGTGGTGGCTTTCTTGACCAA 420
QY 421 ACTCAGCAATCAGTCACTGCTCTGTCACTCAGTGTGACGACCCAGAACATCCAGAGAA 480
DB 421 ACTCAGCAATCAGTCACTGCTCTGTCACTCAGTGTGACGACCCAGAACATCCAGAGAA 480
QY 481 TGTGAGAGGCTGAGAGAGAGAGTGTGAGAGAGAGTGTGAGAGAGAGAGTGTGAGAGAG 540
DB 481 TGTGAGAGGCTGAGAGAGAGAGTGTGAGAGAGAGTGTGAGAGAGAGAGTGTGAGAGAG 540
QY 541 CGGGGAGTGGAGTGTGTTTATGCTCTGAGAAATGCTTGGCTGTGAGGAGAGAGAG 600
DB 541 CGGGGAGTGGAGTGTGTTTATGCTCTGAGAAATGCTTGGCTGTGAGGAGAGAGAG 600
QY 601 CTAGAAAACGAGAACTGCTCTTCTGCTTCTTAAAGAAACAATAAGATCCCTGAATG 660
DB 601 CTAGAAAACGAGAACTGCTCTTCTGCTTCTTAAAGAAACAATAAGATCCCTGAATG 660
QY 661 GACTTTTTTCTAAGGAAAGTGAAGAGCTAAAGTCCACCATCATTTAGAAAGATTTCAT 720
DB 661 GACTTTTTTCTAAGGAAAGTGAAGAGCTAAAGTCCACCATCATTTAGAAAGATTTCAT 720
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121 TGCCCTGTGGCCCGAGGAGCAATGCGCTGCCCATCAACACCCGGTCAAGCTTGAGGT 180  
QY  
181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240  
Db  
181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240  
QY  
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QY  
301 TGCTAAGATCAAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCAACCTCGAAGACATTCT 360  
Db  
301 TGCTAAGATCAAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCAACCTCGAAGACATTCT 360  
QY  
361 GCTCCCCAGTCACAGAGTTCCGGCCCTACATCAGAGAGTGTGCTTCCCTGAGCAA 420  
Db  
361 GCTCCCCAGTCACAGAGTTCCGGCCCTACATCAGAGAGTGTGCTTCCCTGAGCAA 420  
QY  
421 ACTCAGCAATCAGCTCAGCTCCTGTCTCATCAGTGGTGACGACCAAGAACATCCAGAGAA 480  
Db  
421 ACTCAGCAATCAGCTCAGCTCCTGTCTCATCAGTGGTGACGACCAAGAACATCCAGAGAA 480  
QY  
481 TGTGAGAGGCTGAAGGAGACAGTGAAGAAAGCTTGGAGAGCGGAGAGATCAAGGCGAT 540  
Db  
481 TGTGAGAGGCTGAAGGAGACAGTGAAGAAAGCTTGGAGAGCGGAGAGATCAAGGCGAT 540  
QY  
541 CGGGAACTGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAG 600  
Db  
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QY  
601 CTAGAAAACGAAGAACTGCTCCTTCCCTGCTTCTAAGAAAGCAATGAAGTCCCTGAATG 660  
Db  
601 CTAGAAAACGAAGAACTGCTCCTTCCCTGCTTCTAAGAAAGCAATGAAGTCCCTGAATG 660  
QY  
661 GACTTTTTTACTAAGGAAAGTGAGAGCTAAACGTCACCAATCATTAGAATTTTCAAT 720  
Db  
661 GACTTTTTTACTAAGGAAAGTGAGAGCTAAACGTCACCAATCATTAGAATTTTCAAT 720  
QY  
721 GAAACCTGGCTCAGTGAAGAGAAATAGTGTCAAGTGTCCATGATGACGAGAGGTAGA 780  
Db  
721 GAAACCTGGCTCAGTGAAGAGAAATAGTGTCAAGTGTCCATGATGACGAGAGGTAGA 780  
QY  
781 CTTGATAACCAAGAACTTCAATGACAAATATTTTATGTCATTCATTAATGCAACAGAAAA 840  
Db  
781 CTTGATAACCAAGAACTTCAATGACAAATATTTTATGTCATTCATTAATGCAACAGAAAA 840  
QY  
841 AGTATGACTTTTAAAAAATGTTTGAAGGAGGTACCTCTCATTCCTCTAGAGAAAAAG 900  
Db  
841 AGTATGACTTTTAAAAAATGTTTGAAGGAGGTACCTCTCATTCCTCTAGAGAAAAAG 900  
QY  
901 CCTATGTAACCTTCAATTTCCATAACCAATACCTTTATATATATGTAAGTTTATTTAAT 960  
Db  
901 CCTATGTAACCTTCAATTTCCATAACCAATACCTTTATATATATGTAAGTTTATTTAAT 960  
QY  
961 ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAAAATTTCTGATG 1020  
Db  
961 ATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAAAATTTCTGATG 1020  
QY  
1021 TTGATATTTGAGTATAAGCAAAATAATTT 1050  
Db  
1021 TTGATATTTGAGTATAAGCAAAATAATTT 1050

RESULT 10  
US-10-968-432-42  
; Sequence 42, Application US/10968432  
; Publication No. US20050065321A1  
; GENERAL INFORMATION:  
; APPLICANT: Presnell, Scott R.  
; APPLICANT: Xu, Wenfeng  
; APPLICANT: Kindsvogel, Wayne  
; APPLICANT: Chen, Zhi  
; APPLICANT: Hughes, Steven D.

; TITLE OF INVENTION: Human Cytokine Receptor  
; FILE REFERENCE: 01-12C1  
; CURRENT APPLICATION NUMBER: US/10/968,432  
; CURRENT FILING DATE: 2004-10-19  
; PRIOR APPLICATION NUMBER: US 60/279,222  
; PRIOR FILING DATE: 2001-03-27  
; PRIOR APPLICATION NUMBER: US 10/104,919  
; PRIOR FILING DATE: 2002-03-22  
; NUMBER OF SEQ ID NOS: 62  
; SOFTWARE: Fast-Seq for Windows Version 3.0  
; SEQ ID NO 42  
; LENGTH: 1050  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (5)...(589)  
US-10-968-432-42  
  
Query Match 94.4%; Score 1048.4; DB 9; Length 1050;  
Best Local Similarity 99.9%; Pred. No. 2.2e-215;  
Matches 1049; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
  
QY 1 AACAGGCTCTCCTCTCAGTTTATCAACTTTTGACACTTTGCGATCGGTGATGGCTGTCT 60  
Db 1 AACAGGCTCTCCTCTCAGTTTATCAACTTTTGACACTTTGCGATCGGTGATGGCTGTCT 60  
QY 61 GCAGAAATCTATCAGTTTTCCTTATGGGACATTTGGCCCGCAGCTGCTTCTCAT 120  
Db 61 GCAGAAATCTATCAGTTTTCCTTATGGGACATTTGGCCCGCAGCTGCTTCTCAT 120  
QY 121 TGCCCTGTGGCCCGAGGAGCAATGCGTGCCTCAACACCCGGTGAAGCTTGAGGT 180  
Db 121 TGCCCTGTGGCCCGAGGAGCAATGCGTGCCTCAACACCCGGTGAAGCTTGAGGT 180  
QY 181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240  
Db 181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGCCAG 240  
QY 241 CCTTGAGATAAACACACAGAGCTCCGGCTCATCGGGGAGAAACTGTTCCGAGAGTCCAG 300  
Db 241 CCTTGAGATAAACACACAGAGCTCCGGCTCATCGGGGAGAAACTGTTCCGAGAGTCCAG 300  
QY 301 TGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCACTTCACTTCCCTGAGACATTCT 360  
Db 301 TGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCACTTCACTTCCCTGAGACATTCT 360  
QY 361 GCTCCCCCAGTCACAGAGTTCCGGCCCTACATGACGAGGTGGTGGCTTTTCTTGACCAA 420  
Db 361 GCTCCCCCAGTCACAGAGTTCCGGCCCTACATGACGAGGTGGTGGCTTTTCTTGACCAA 420  
QY 421 ACTCAGCAATCAGCTCAGCTCCTGTCTCATCAGTGGTGACGACCAAGAACATCCAGAGAA 480  
Db 421 ACTCAGCAATCAGCTCAGCTCCTGTCTCATCAGTGGTGACGACCAAGAACATCCAGAGAA 480  
QY 481 TGTGAGAGGCTGAAGGAGACAGTGAAGAAAGCTTGGAGAGCGGAGAGATCAAGGCGAT 540  
Db 481 TGTGAGAGGCTGAAGGAGACAGTGAAGAAAGCTTGGAGAGCGGAGAGATCAAGGCGAT 540  
QY 541 CGGGAACTGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAG 600  
Db 541 CGGGAACTGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAG 600  
QY 601 CTAGAAAACGAAGAACTGCTCCTTCCCTGCTTCTAAGAAAGCAATGAAGTCCCTGAATG 660  
Db 601 CTAGAAAACGAAGAACTGCTCCTTCCCTGCTTCTAAGAAAGCAATGAAGTCCCTGAATG 660  
QY 661 GACTTTTTTACTAAGGAAAGTGAGAGCTAAACGTCACCAATCATTAGAATTTTCAAT 720  
Db 661 GACTTTTTTACTAAGGAAAGTGAGAGCTAAACGTCACCAATCATTAGAATTTTCAAT 720  
QY 721 GAAACCTGGCTCAGTGAAGAGAAATAGTGTCAAGTGTCCATGATGACGAGAGGTAGA 780  
Db 721 GAAACCTGGCTCAGTGAAGAGAAATAGTGTCAAGTGTCCATGATGACGAGAGGTAGA 780



Db 721 GAAACCTGGCTCAGTTGAAGAGAAATAGTGTCAAGTTGTCATGAGACGAGGTAGA 780  
Qy 781 CTTGATAACCAAGATTCATTGACAAATATTTATTTGTCATTGATTAATGCAACAGAAAA 840  
Db 781 CTTGATAACCAAGATTCATTGACAAATATTTATTTGTCATTGATTAATGCAACAGAAAA 840  
Qy 841 AGTATGTACTTTAAAAATTTGTTGAAGAGGTACCTCTCATTTCTCTAGAGAAAAAG 900  
Db 841 AGTATGTACTTTAAAAATTTGTTGAAGAGGTACCTCTCATTTCTCTAGAGAAAAAG 900  
Qy 901 CCTATGTACTTTCCATATCCATACCAATACCTTTATATATGATGATTTATTTATTAAGT 960  
Db 901 CCTATGTACTTTCCATATCCATACCAATACCTTTATATATGATGATTTATTTATTAAGT 960  
Qy 961 ATACATTTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAAAATTTATCTGATG 1020  
Db 961 ATACATTTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAAAATTTATCTGATG 1020  
Qy 1021 TTGATATTTGAGTATTAAGCAAAATATATT 1050  
Db 1021 TTGATATTTGAGTATTAAGCAAAATATATT 1050

## RESULT 11

US-11-045-944-40  
; Sequence 40, Application US/11045944  
; Publication No. US20050124796A1  
; GENERAL INFORMATION:  
; APPLICANT: Preenell, Scott R.  
; APPLICANT: Xu, Wenfeng  
; APPLICANT: Kindsvogel, Wayne  
; APPLICANT: Chen, Zhi  
; TITLE OF INVENTION: Mouse Cytokine Receptor  
; FILE REFERENCE: 01-08D1  
; CURRENT APPLICATION NUMBER: US/11/045,944  
; CURRENT FILING DATE: 2005-01-28  
; PRIOR APPLICATION NUMBER: US 60/273,035  
; PRIOR FILING DATE: 2001-03-02  
; PRIOR APPLICATION NUMBER: US 60/279,232  
; PRIOR FILING DATE: 2001-03-27  
; PRIOR APPLICATION NUMBER: US 10/090,365  
; PRIOR FILING DATE: 2002-03-04  
; NUMBER OF SEQ ID NOS: 49  
; SOFTWARE: FastSeq for Windows Version 3.0  
; SEQ ID NO 40  
; LENGTH: 1050  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (50)...(589)  
US-11-045-944-40

Query Match 94.4%; Score 1048.4; DB 10; Length 1050;

Best Local Similarity 99.9%; Pred. No. 2.2e-215;

Matches 1049; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 AACAGGCTCTCCTCAGTTTATCAACTTTTGACACTTTGTCGGATCGGTGATGCGTGCCT 60  
Db 1 AACAGGCTCTCCTCAGTTTATCAACTTTTGACACTTTGTCGGATCGGTGATGCGTGCCT 60  
Qy 61 GCAGAAATCTATGAGTTTTCCTTATGCGGACTTTTGGCCGCCAGCTGCTGCTTCTCAT 120  
Db 61 GCAGAAATCTATGAGTTTTCCTTATGCGGACTTTTGGCCGCCAGCTGCTGCTTCTCAT 120  
Qy 121 TGGCCTGTGGCCAGAGGCAATGCGTGCCTATCAACCCGGTGCAGGTTGAGGT 180  
Db 121 TGGCCTGTGGCCAGAGGCAATGCGTGCCTATCAACCCGGTGCAGGTTGAGGT 180  
Qy 181 GTCCAACTTCAGAGCCGCTACATCGTCAACCCGACCTTTATGCTGCGCAAGGCGCAG 240  
Db 181 GTCCAACTTCAGAGCCGCTACATCGTCAACCCGACCTTTATGCTGCGCAAGGCGCAG 240

Qy 241 CTTTGAGATTAACAAACACAGACGTCGGCTCATCGGGAGAAACTGTTCCGAGAGTCAG 300  
Db 241 CTTTGAGATTAACAAACACAGACGTCGGCTCATCGGGAGAAACTGTTCCGAGAGTCAG 300  
Qy 301 TGCTRAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCCACCTCGAAGACATTCT 360  
Db 301 TGCTRAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCCACCTCGAAGACATTCT 360  
Qy 361 GCTTCCCCCACTCAGACAGGTTCGGGCCCTACATGACGAGAGTGGTGCCTTTCTCGACCA 420  
Db 361 GCTTCCCCCACTCAGACAGGTTCGGGCCCTACATGACGAGAGTGGTGCCTTTCTCGACCA 420  
Qy 421 ACTCAGCAATCAGCTCAGCTCCTGTCACATCAGTGTGAGCAGCAGAAATCCAGAGAA 480  
Db 421 ACTCAGCAATCAGCTCAGCTCCTGTCACATCAGTGTGAGCAGCAGAAATCCAGAGAA 480  
Qy 481 TGTCAAGAGCTGAAGGAGACAGTGAAGAAAGCTTTGAGAGAGCGGAGAGATCAAGCGAT 540  
Db 481 TGTCAAGAGCTGAAGGAGACAGTGAAGAAAGCTTTGAGAGAGCGGAGAGATCAAGCGAT 540  
Qy 541 CGGGAACTGGACCTGCTGTTATGTCCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAG 600  
Db 541 CGGGAACTGGACCTGCTGTTATGTCCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAG 600  
Qy 601 CTAGAAAAAGAGAACTGCTCTTCTGCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 660  
Db 601 CTAGAAAAAGAGAACTGCTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 660  
Qy 661 GACTTTTTTACTTAAAGGAAAGTGAAGAGCTTAAGCTCCACCATCATTTAGAGATTTTCACAT 720  
Db 661 GACTTTTTTACTTAAAGGAAAGTGAAGAGCTTAAGCTCCACCATCATTTAGAGATTTTCACAT 720  
Qy 721 GAAACCTGGCTCAGTTGAAAGAGAAATAGTGTCAAGTTGTCCATGAGACGAGAGGTAGA 780  
Db 721 GAAACCTGGCTCAGTTGAAAGAGAAATAGTGTCAAGTTGTCCATGAGACGAGAGGTAGA 780  
Qy 781 CTTGATAACCAAGATTCATTGACAAATATTTATTTGTCATTGATTAATGCAACAGAAAA 840  
Db 781 CTTGATAACCAAGATTCATTGACAAATATTTATTTGTCATTGATTAATGCAACAGAAAA 840  
Qy 841 AGTATGTACTTTAAAAATTTGTTGAAGAGGTACCTCTCATTTCTCTAGAGAAAAAG 900  
Db 841 AGTATGTACTTTAAAAATTTGTTGAAGAGGTACCTCTCATTTCTCTAGAGAAAAAG 900  
Qy 901 CCTATGTACTTTCCATATCCATACCAATACCTTTATATATGATGATTTATTTATTAAGT 960  
Db 901 CCTATGTACTTTCCATATCCATACCAATACCTTTATATATGATGATTTATTTATTAAGT 960  
Qy 961 ATACATTTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAAAATTTATCTGATG 1020  
Db 961 ATACATTTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAAAATTTATCTGATG 1020  
Qy 1021 TTGATATTTGAGTATTAAGCAAAATATATT 1050  
Db 1021 TTGATATTTGAGTATTAAGCAAAATATATT 1050

## RESULT 12

US-09-751-797-7

; Sequence 7, Application US/09751797

; Patent No. US20010024652A1

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Louhed, Jamila

; APPLICANT: Renaud, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; FILE REFERENCE: (Tifs) The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: IUD 5543.2

; CURRENT APPLICATION NUMBER: US/09/751,797

; CURRENT FILING DATE: 2000-12-29

; PRIOR APPLICATION NUMBER: 09/419,568

; PRIOR FILING DATE: 1999-10-18

; PRIOR APPLICATION NUMBER: US09/178,973



; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 7
; LENGTH: 1119
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-09-751-797-7

Query Match 94.3%; Score 1047.8; DB 3; Length 1119;
Best Local Similarity 97.0%; Pred. No. 3e-215;
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;

Qy 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGGGATCGGTGATGGCTGTCCT 60
Db 3 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGGGATCGGTGATGGCTGTCCT 62

Qy 61 GCAGAAATCTATGAGTTTTTCCCTTATGGGACATTTTGGCGCCACAGCTGCCTTCTCAT 120
Db 63 GCAGAAATCTATGAGTTTTTCCCTTATGGGACATTTTGGCGCCACAGCTGCCTTCTCAT 122

Qy 121 TGCCTCTGGGCCCAGAGGCAAAATGCGTGCCCATCAACACCCGGTGCAAGCTTGAGGT 180
Db 123 TGCCTCTGGGCCCAGAGGCAAAATGCGTGCCCGTCAACACCCGGTGCAAGCTTGAGGT 182

Qy 181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGCCAG 240
Db 183 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGCCAG 242

Qy 241 CTTTGCAGATAACACAGAGTCCGGCTCATCGGAGGAGAACTGTTCCGAGAGTCA 300
Db 243 CTTTGCAGATAACACAGAGTCCGGCTCATCGGAGGAGAACTGTTCCGAGAGTCA 302

Qy 301 TGCTAAGGATCAGTGCTACCTGATGAAGAGGCTGCTCAACTTCACTCCCTGGAAGACATCT 360
Db 303 TGCTAAGGATCAGTGCTACCTGATGAAGAGGCTGCTCAACTTCACTCCCTGGAAGACATCT 362

Qy 361 GCTCCCCAGTCACAGAGTTCGGCCCTCATCGAGGAGTGTGCTTCTTCGACCA 420
Db 363 GCTCCCCAGTCACAGAGTTCGGCCCTCATCGAGGAGTGTGCTTCTTCGACCA 422

Qy 421 ACTCAGCAATCAGTCTCAGTCTCTGTCAATCAGTGGTGACGACCAAGAACATCCAGAGAA 480
Db 423 ACTCAGCAATCAGTCTCAGTCTCTGTCAATCAGTGGTGACGACCAAGAACATCCAGAGAA 482

Qy 481 TGTCAAGGCTGAAGGAGACAGTGAAAGCTTTGGAGAGCGGAGAGATCAAGGCGAT 540
Db 483 TGTCAAGGCTGAAGGAGACAGTGAAAGCTTTGGAGAGCGGAGAGATCAAGGCGAT 542

Qy 541 CGGGAACTGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAG 600
Db 543 TGGGAACTGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAG 602

Qy 601 CTAGAAACGAGAACTGCTCTCTGCTGCTTTCTAAAAGAAACAATAAGATCCCTGAATG 660
Db 603 CTAGAAACGAGAACTGCTCTCTGCTGCTTTCTAAAAGAAACAATAAGATCCCTGAATG 662

Qy 661 GACTTTTTTCTAAAGGAAAGTGAGAGCTCAACGTCACCATCATATTAGAGATTTCAT 720
Db 663 GACTTTTTTCTAAAGGAAAGTGAGAGCTCAACGTCACCATCATATTAGAGATTTCAT 722

Qy 721 GAAACCTGGCTCAGTTGAAAGGAAATAGTGTCAAGTTGTCATGTGACCCAGAGGTAGA 780
Db 723 GAAACCTGGCTCAGTTGAAAGGAAATAGTGTCAAGTTGTCATGTGACCCAGAGGTAGA 782

Qy 781 CTTGATAACCAAGAGATTTCATTGACATAATTTTATTGTCATGATGATGCAACAGAGAA 840
Db 783 CTTGATAACCAAGAGATTTCATTGACATAATTTTATTGTCATGATGATGCAACAGAGAA 842

Qy 841 AGTATGTAATTTAAAGAAATGTTTGAAGAGGTTTACCTCTCATTTCTCTAGAGAGAAAG 900
Db 843 ATAATGTAATTTAAAGAAATGTTTGAAGAGGTTTACCTCTCATTTCTCTTAGAGAGAAAG 902

RESULT 13
US-10-627-273-7
; Sequence 7, Application US/10627273
; Publication No. US20040110189A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: (TIFF) The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.2
; CURRENT APPLICATION NUMBER: US/10/627,273
; CURRENT FILING DATE: 2003-07-25
; PRIOR APPLICATION NUMBER: US/09/751,797
; PRIOR FILING DATE: 2000-12-29
; PRIOR APPLICATION NUMBER: 09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 7
; LENGTH: 1119
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-10-627-273-7

Query Match 94.3%; Score 1047.8; DB 7; Length 1119;
Best Local Similarity 97.0%; Pred. No. 3e-215;
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;

Qy 1 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCGGTGATGGCTGTCCT 60
Db 3 AACAGGCTCTCTCTCAGTTATCAACTTTTGACACTTTGTGCGATCTCTGATGGCTGTCCT 62

Qy 61 GCAGAAATCTATGAGTTTTTCCCTTATGGGACATTTGGCGCCACAGTGCCTGCTTCAT 120
Db 63 GCAGAAATCTATGAGTTTTTCCCTTATGGGACATTTGGCGCCACAGTGCCTGCTTCAT 122

Qy 121 TGCCTCTGGGCCCAGAGGCAAAATGCGTCCCATCAACACCCGGTGCAAGCTTGAGGT 180
Db 123 TGCCTCTGGGCCCAGAGGCAAAATGCGTCCCATCAACACCCGGTGCAAGCTTGAGGT 182

Qy 181 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGCCAG 240
Db 183 GTCCAACTTCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGCCAG 242

Qy 241 CTTTGCAGATAACACAGAGTCCGGCTCATCGGAGGAGAACTGTTCCGAGAGTCA 300
Db 243 CTTTGCAGATAACACAGAGTCCGGCTCATCGGAGGAGAACTGTTCCGAGAGTCA 302

Qy 301 TGCTAAGGATCAGTGCTTACCTGATGAAGAGGCTGCTCAACTTCACTCCCTGGAAGACATCT 360
Db 303 TGCTAAGGATCAGTGCTTACCTGATGAAGAGGCTGCTCAACTTCACTCCCTGGAAGAGTCT 362

Qy 361 GCTCCCCAGTCACAGAGTTCGGCCCTCATCGAGGAGTGTGCTTCTTCGACCA 420

	Db	363	GCTCCCCGAGTCAGACAGGTTCAGCGCCTCATCATGCAGGAGTGGTATCTTTCCCTGACCNA	422
	Qy	421	ACTCAGCAATCAGCTCAGCTCCTGTGCATCATCAGTGGTGACGACACAGAACAATCCAGAAGAA	480
	Db	423	ACTCAGCAATCAGCTCAGCTCCTGTGCATCATCAGCGGTGACGACAGAACATCCAGAAGNA	482
	Qy	481	TGTCAGAAGGCTGAAGGAGACAGTGA AAAAGCTTTGGAGAGAGCGGAGAGATCAAAGCGAT	540
	Db	483	TGTCAGAAGGCTGAAGGAGACAGTGA AAAAGCTTTGGAGAGAGTGGAGAGATCAAGGCGAT	542
	Qy	541	CGGGGAACCTGCACCTGCTGTTTTATGTCTCTGAGAAATGCTTTGGCTCTGACGAGAGAAG	600
	Db	543	TGGGGAACTGGA C CTGCTGTTTTATGTCTCTGAGAAATGCTTTGGCTCTGACGAGAGAAG	602
	Qy	601	CTAGAAAACGAAGA ACTGCTCCTTCTCGCCCTCTTAAAAAGAACATAAGATCCCTGTAATG	660
	Db	603	CTAGAAAACGAAGA ACTGCTCCTTCTCGCCCTCTTAAAAAGAACATAAGATCCCTGTAATG	662
	Qy	661	GACTTTTTTACTAAAGAAAGTGAGANGCTTAA CGTCCACCATCATTTAGAAGATTTCACAT	720
	Db	663	GACTTTTTTACTAAAGAAAGTGAGANGCTTAA CGTCCATCATCATTTAGAAGATTTCACAT	722
	Qy	721	GA AACCTGGCTCAGTTGAAAAGAGAAAATAGTGTCAAGTTGTCCATGAGACCAGAGGTAGA	780
	Db	723	GA AACCTGGCTCAGTTGAAAAGAGAAAATAGTGTCAAGTTGTCCATGAGACCAGAGGTAGA	782
	Qy	781	CTTGATTAACCA AAGATTTCATGACAATATTTTATTTGTCA TTGATTAATGCAACAGAAAA	840
	Db	783	CTTGATTAACCA AAGATTTCATGACAATATTTTATTTGTCA TTGATTAATGCAACAGAAAA	842
	Qy	841	AGTATGTACTTTTAAAAATTTGTTTGAAGGAGGTTACCTCTCATTCCTCTTAGAAGAAAAG	900
	Db	843	ATAATGTACTTTTAAAAATTTGTTTGAAGGAGGTTACCTCTCATTCCTTTAGAAAAAAG	902
	Qy	901	CCTATGTAACCTTCATTTCCATAACCA TACTTTATATATATGATGAAGTTTATTTATTAAGT	960
	Db	903	CTTATGTAACTTCATTTCCATA TCCAATA TTTTATATATGATGAAGTTTATTTATTAAGT	962
	Qy	961	ATACATTTTATTTATGTCA GTTTTATTAATGGA TTTTATTTATAGAAAAATTTATCTGATG	1020
	Db	963	ATACATTTTATTTATGTCA GTTTTATTAATGGA TTTTATTTATAGAAAAATTTATCTGCTA	1022
	Qy	1021	TTGATATTTTCAGTATAAGCAATAATATTTTATGATAATAACTATAGAAA CAAGATATCT	1080
	Db	1023	TTGATATTT - AGTATAAGGCCAATAATATTTTATGACAA TAACATATGAAA CAAGATATCT	1081
	Qy	1081	TAGCCTTTAATAAACACATGA ATCATATAA 1111	
	Db	1082	TAGCCTTTAATAAACACATGA ATCATATAA 1112	

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RESULT 14
US-09-746-375-37
; Sequence 37, Application US/09746375
; Publication No. US20030170823A1
; GENERAL INFORMATION:
; APPLICANT: Preenell, Scott R.
; APPLICANT: Kindsvogel, Wayne
; TITLE OF INVENTION: NOVEL CYTOKINE ZCYTO18
; FILE REFERENCE: 99-106
; CURRENT APPLICATION NUMBER: US/09/746,375
; CURRENT FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 60/172,105
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: US 60/****,***
; PRIOR FILING DATE: 2000-12-01
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 37
; LENGTH: 778
; TYPE: DNA
; ORGANISM: mus musculus

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; FEATURE:
; NAME/KEY: CDS
; LOCATION: (47) ... (583)
US-09-746-375-37

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Best Local Similarity 98.3%; Pred. No. 9.5e-153;
Matches 765; Conservative 0; Mismatches 13; Indels 0; Gaps 0;

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Db  1  AGGCTCTCCTCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGCTGCTCTCATGCA 60
    |
Qy  64  GAAATCATAGATTTTCCCTTATGCGGACTTTGGCGCCAGCTGCTGCTTCTCATGTC 123
    |
Db  61  GAAATCATAGATTTTCCCTTATGCGGACTTTGGCGCCAGCTGCTGCTTCTCATGTC 120
    |
Qy  124  CTTGTGGGCCAGGAGGCAATGCGCTGCCCATCAACCCCGTGCAAGCTTCAGAGTGC 183
    |
Db  121  CTTGTGGGCCAGGAGGCAATGCGCTGCCCGTCAACCCCGTGCAAGCTTCAGAGTGC 180
    |
Qy  184  CAATCTCCAGCAGCCGTACATCGTCAACCGCACCTTTTATGCTGGCCAAAGAGGCCAGCCT 243
    |
Db  181  CAATCTCCAGCAGCCGTACATCGTCAACCGCACCTTTTATGCTGGCCAAAGAGGCCAGCCT 240
    |
Qy  244  TGCAGATAACAAACACAGAGCTCCGGCTCATTCGGGGAGAACTCTTCCGAGGAGTCAAGTGC 303
    |
Db  241  TGCAGATAACAAACACAGAGTCCGGCTCATTCGGGGAGAACTCTTCCGAGGAGTCAAGTGC 300
    |
Qy  304  TAAGGATCAGTGCTACCTGATGAAGCAGGTGCTCAACTTTCACCCCTGGAGACATTCCTGCT 363
    |
Db  301  TAAGGATCAGTGCTACCTGATGAAGCAGGTGCTCAACTTTCACCCCTGGAGACATTCCTGCT 360
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Qy  364  CCCCCAGTCAGACAGGTTCCGGCCCTACATGACGAGGAGTGGTCCCTTCTCTGACCCAAACT 423
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Qy  424  CAGCAATCAGCTCAGTCTCCTGTCAATCATAGTGGTGCAGCCAGAACATCCAGAAGAATGT 483
    |
Db  421  CAGCAATCAGCTCAGTCTCCTGTCAATCATAGCAGGTGCAGCCAGAACATCCAGAAGAATGT 480
    |
Qy  484  CAGAAGGCTGAAGGAGACAGTGAAGGCTTGAGAGAGCGGAGAGATCAAGCGATCGG 543
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Qy  544  GGAACCTGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGACGAGAGAGAGCTA 603
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Qy  604  GAAACGGAAGAACTGCTCTCTTCTGCTCTTCTAAAAAGAACATAAGATCCCTGAATGGAC 663
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Db  601  GAAACGGAAGAACTGCTCTCTTCTGCTCTTCTAAAAAGAACATAAGATCCCTGAATGGAC 660
    |
Qy  664  TTTTCTTAAAGGAAAGTGAGAGCTAACGTCACCATCATTTAGAAGATTTTCAATGAA 723
    |
Db  661  TTTTCTTAAAGGAAAGTGAGAGCTAACGTCATCATTTAGAAGATTTTCAATGAA 720
    |
Qy  724  ACCTGGCTCAGTTGAAGAGAAATAGTGTCAAGTTGTCCATGAGACCGAGAGTAC 781
    |
Db  721  ACCTGGCTCAGTTGAAGAGAAATAGTGTCAAGTTGTCCATGAGACCGAGAGTAC 778
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RESULT 15
US-10-395-741B-37
; Sequence 37, Application US/10395741B
; Publication No. US20040023341A1
; GENERAL INFORMATION:
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Hughes, Steven D.
; APPLICANT: Chandrasekhar, Yasmin A.
; TITLE OF INVENTION: ANTI-IL-TTF ANTIBODIES AND METHODS OF
; TITLE OF INVENTION: USING IN INFLAMMATION

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Job time : 558.632 secs

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; CURRENT FILING DATE: 2003-03-24  
; PRIOR APPLICATION NUMBER: US 60/366,842  
; PRIOR FILING DATE: 2002-03-22  
; NUMBER OF SEQ ID NOS: 67  
; SOFTWARE: FastSeq for Windows Version 3.0  
; SEQ ID NO 37  
; LENGTH: 778  
; TYPE: DNA  
; ORGANISM: mus musculus  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (47)...(583)  
US-10-395-741B-37

Query Match 68.2%; Score 757.2; DB 7; Length 778;  
Best Local Similarity 98.3%; Pred. No. 9.5e-153;  
Matches 765; Conservative 0; Mismatches 13; Indels 0; Gaps 0;  
  
Qy 4 AGGCTCTCCTCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGGCTGTCTCTGCA 63  
Db |||||||  
Qy 1 AGGCTCTCCTCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGGCTGTCTCTGCA 60  
Db |||||||  
  
Qy 64 GAATCTATGAGTTTTCCTTTATGGGACTTTTGGCCGCCAGCTGCCCTTCTCTCATTTGC 123  
Db |||||||  
Qy 61 GAATCTATGAGTTTTCCTTTATGGGACTTTTGGCCGCCAGCTGCCCTTCTCTCATTTGC 120  
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Qy 124 CCTGTGGGCCCAGGAGCAATGCGTCCCATCAACACCCGGTGCAAGCTTGAGGTGTC 183  
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Qy 181 CAACCTTCAGAGCGGTACATCGTCAACCGCACCTTTATGTGCGCAAGAGGCCAGCCT 240  
Db |||||||  
  
Qy 244 TGCAGATAACAACACAGAGTCCGGCTCATCGGGGAAACTGTTCCGAGAGTCAGTGC 303  
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Qy 241 TGCAGATAACAACAGAGTCCGGCTCATCGGGGAAACTGTTCCGAGAGTCAATGC 300  
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Qy 304 TAAGGATCAGTGTACCTGTATGAAGCAGGTGCTCAACTTTCACCTTGAAGACATTTCTGCT 363  
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Qy 364 CCCCAGTCAGACAGGTTCCGGCCCTACATGACGAGAGGTGTGCTTTCTTGACCAAACT 423  
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Qy 361 CCCCAGTCAGACAGGTTCCAGCCCTACATGACGAGAGGTGTGCTTTCTTGACCAAACT 420  
Db |||||||  
  
Qy 424 CAGCAATCAGCTCAGCTCCTGTACATCAGTGTGACGACCAGAACATCCAGAGAATGT 483  
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Qy 421 CAGCAATCAGCTCAGCTCCTGTACATCAGCGGTGACGACCAGAACATCCAGAGAATGT 480  
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Qy 484 CAGAAGCTGAAGGAGACAGTGAAAGCTTTGGAGAGACGGGAGAGATCAAGCGATCGG 543  
Db |||||||  
Qy 481 CAGAAGCTGAAGGAGACAGTGAAAGCTTTGGAGAGAGTGGAGAGATCAAGCGATGG 540  
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Qy 544 GGAAGTGAACCTGTGTTTATGTCTTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTA 603  
Db |||||||  
Qy 541 GGAAGTGAACCTGTGTTTATGTCTTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTA 600  
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Qy 664 TTTTCTTAAAGGAAAGTGAAGAGCTTAACGTCACCATCATTTAGAGATTTTCATGAA 723  
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Qy 724 ACCTGGCTCAGTTGAAAGAGAAAAATAGTGTCAAGTTGTTCATGTAGACCCAGAGGTAGAC 781  
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Qy 721 ACCTGGCTCAGTTGAAAGAGAAAAATAGTGTCAAGTTGTTCATGTAGACCCAGAGGTAGAC 778  
Db |||||||

GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

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Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 4168288 seqs, 259477437 residues

Total number of hits satisfying chosen parameters: 8336576

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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Published Applications NA New.\*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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4	555.2	50.0	7445	7	US-11-177-987-8
5	525.6	47.3	1152	7	US-11-102-240-153
6	407.6	36.7	680	7	US-11-177-987-25
7	217.2	19.5	418	7	US-11-177-987-18
8	127.6	11.5	4797	7	US-11-177-987-26
9	61	5.5	161874	7	US-11-121-086-75
10	57.6	5.2	151169	7	US-11-121-086-38
11	55.6	5.0	171486	7	US-11-121-086-105
12	55.6	5.0	394468	6	US-10-995-561-13473
13	55.2	5.0	26772	6	US-10-995-561-13313
14	55.2	5.0	54946	6	US-10-995-561-13479
15	54.6	4.9	171486	7	US-11-121-086-105
16	54.4	4.9	394468	6	US-10-995-561-13473
17	53.2	4.8	171602	7	US-11-121-086-25
18	53	4.8	181172	7	US-11-121-086-41
19	52.8	4.8	105550	6	US-10-995-561-13235
20	51.8	4.7	173602	7	US-11-121-086-25
21	51.6	4.6	146656	7	US-11-121-086-68
22	51.6	4.6	151169	7	US-11-121-086-38
23	51	4.6	20317	6	US-10-995-561-13460

C	24	50.2	4.5	147700	6	US-10-857-780-3	Sequence 3, Appli
	25	50	4.5	119036	6	US-10-995-561-13314	Sequence 13314, A
C	26	49.8	4.5	7720	6	US-10-515-481-10	Sequence 10, Appl
	27	49.6	4.5	161874	7	US-11-121-086-75	Sequence 75, Appl
	28	49.4	4.4	12872	6	US-10-995-561-13511	Sequence 13511, A
C	29	49.2	4.4	201990	6	US-10-995-561-13303	Sequence 13303, A
	30	49.2	4.4	398287	6	US-10-995-561-13396	Sequence 13396, A
C	31	49	4.4	175416	7	US-11-121-086-43	Sequence 43, Appl
	32	48.8	4.4	14941	6	US-10-821-234-771	Sequence 771, App
	33	48.6	4.4	47444	6	US-10-995-561-13354	Sequence 13354, A
C	34	48.6	4.4	105550	6	US-10-995-561-13235	Sequence 13235, A
	35	48.4	4.4	1483	6	US-10-750-185-39052	Sequence 39052, A
	36	48.4	4.4	49979	6	US-10-995-561-13443	Sequence 13443, A
C	37	48.2	4.3	119036	6	US-10-995-561-13314	Sequence 13314, A
	38	48.2	4.3	134499	7	US-11-117-187-192	Sequence 192, App
	39	48.2	4.3	182314	7	US-11-112-908-45	Sequence 45, Appl
C	40	48.2	4.3	184868	7	US-11-121-086-88	Sequence 88, Appl
	41	48.2	4.3	201990	6	US-10-995-561-13303	Sequence 13303, A
C	42	48	4.3	201	6	US-10-995-561-74696	Sequence 74696, A
	43	48	4.3	197781	7	US-11-112-908-34	Sequence 34, Appl
C	44	47.8	4.3	1483	6	US-10-750-185-39052	Sequence 39052, A
	45	47.6	4.3	7720	6	US-10-515-481-10	Sequence 10, Appl

ALIGNMENTS

RESULT 1

US-11-177-987-9  
; Sequence 9, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fact  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; PRIOR FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-11-177-987-9

Query Match	100.0%	Score 1111;	DB 7;	Length 1111;
Best Local Similarity	100.0%;	Pred. No. 1e-250;		
Matches 1111;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
QY	1	AACAGGCTCTCTCTCAGTATCAACTTTTGACATCTTGCGCATCGGTGATGCTGCTCT	60	
DB	1	AACAGGCTCTCTCTCAGTATCAACTTTTGACATCTTGCGCATCGGTGATGCTGCTCT	60	
QY	61	GCAGAAATCTATGAGTTTTTCCTTATGGGACTTTGGCGCGCAGTCTGCTTCTCAT	120	
DB	61	GCAGAAATCTATGAGTTTTTCCTTATGGGACTTTGGCGCGCAGTCTGCTTCTCAT	120	
QY	121	TGCGCTGTGGGCGGAGGCAAAATCGCTTGCCCATCAACCCGTCGCAAGTTGAGGT	180	
DB	121	TGCGCTGTGGGCGGAGGCAAAATCGCTTGCCCATCAACCCGTCGCAAGTTGAGGT	180	
QY	181	GTCCAACTTCCAGCAGCCGCTACATCGTCAACCGCACCTTTATGCTGGCCAAAGAGGCCAG	240	

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Db 181 GTCAAATCTCCAGCAGCGGTACATCGTCAACCGCACCTTTATGTGTCGCAAGGAGCCAG 240
Qy 241 CCTTGCAGATAACAAACAGACGCTCCGGCTCATCGGGGAGAACTGTTCGAGAGTCAG 300
Db 241 CCTTGCAGATAACAAACAGACGCTCCGGCTCATCGGGGAGAACTGTTCGAGAGTCAG 300
Qy 301 TGCTAAGGATCAGTGTCTACCTGTATGAACAGAGTCTCAACTTTCACCCCTGGAAGACATTC 360
Db 301 TGCTAAGGATCAGTGTCTACCTGTATGAACAGAGTCTCAACTTTCACCCCTGGAAGACATTC 360
Qy 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCGCAGAGGTGTGCTTCTCTGACCAA 420
Db 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCGCAGAGGTGTGCTTCTCTGACCAA 420
Qy 421 ACTCAGCAATCAGCTCAGCTCCTGTCTACATCAGTGTGACGACAGCAACATCCAGAGAA 480
Db 421 ACTCAGCAATCAGCTCAGCTCCTGTCTACATCAGTGTGACGACAGCAACATCCAGAGAA 480
Qy 481 TGTGAGAAAGCTGAAGGAGACAGTGAAAGAACTTTGGAGAGCGGAGAGATCAAGCGAT 540
Db 481 TGTGAGAAAGCTGAAGGAGACAGTGAAAGAACTTTGGAGAGCGGAGAGATCAAGCGAT 540
Qy 541 CGGGGAATCGAATCTGCTGTTTATGTCTCTGAGAAATGCTTGCCTGTGAGCGAGAGAG 600
Db 541 CGGGGAATCGAATCTGCTGTTTATGTCTCTGAGAAATGCTTGCCTGTGAGCGAGAGAG 600
Qy 601 CTAGAAACGAGAACTGCTCCTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 660
Db 601 CTAGAAACGAGAACTGCTCCTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 660
Qy 720 GACTTTTTTACTAAGGAAAGTGAAGAGCTAACTGCTCAACATCAATTAGAGATTTTCACAT 720
Db 720 GACTTTTTTACTAAGGAAAGTGAAGAGCTAACTGCTCAACATCAATTAGAGATTTTCACAT 720
Qy 780 GAAACCTGGCTCAGTGTGAAAGAGAAATAGTGTCAAGTTGTCCATGAGACCAAGAGGTAGA 780
Db 780 GAAACCTGGCTCAGTGTGAAAGAGAAATAGTGTCAAGTTGTCCATGAGACCAAGAGGTAGA 780
Qy 840 CTTCATACCAACAGATTCATTCAGCAATATTTTGTCAATTCATGATCAATGATGATGATG 840
Db 840 CTTCATACCAACAGATTCATTCAGCAATATTTTGTCAATTCATGATCAATGATGATGATG 840
Qy 900 AGTATGTACTTTTAAAAAATTTTGTGAAAGAGGTTACCTCTCATCTCTAGAGAGAAAG 900
Db 900 AGTATGTACTTTTAAAAAATTTTGTGAAAGAGGTTACCTCTCATCTCTAGAGAGAAAG 900
Qy 960 CCTATGTAATTCATTTCCATAAACATACTTTATATATATATATATATATATATATATAT 960
Db 960 CCTATGTAATTCATTTCCATAAACATACTTTATATATATATATATATATATATATATAT 960
Qy 1020 ATACATTTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAAATTTATCTGATG 1020
Db 1020 ATACATTTTATTTATGTCAGTTTATTAATATGATTTATTTATAGAAAATTTATCTGATG 1020
Qy 1080 TTGATATTTAGTATATAAGCAAAATAATTTATGATATAATACTATAGAAAATTTATCT 1080
Db 1080 TTGATATTTAGTATATAAGCAAAATAATTTATGATATAATACTATAGAAAATTTATCT 1080
Qy 1111 TAGGCTTTAATAACACATGAATATCATAAA 1111
Db 1111 TAGGCTTTAATAACACATGAATATCATAAA 1111
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RESULT 2  
US-11-177-987-7  
; Sequence 7, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664

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; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-11-177-987-7  
  
Query Match 94.3%; Score 1047.8; DB 7; Length 1119;  
Best Local Similarity 97.0%; Pred. No. 6.4e-236;  
Matches 1078; Conservative 0; Mismatches 32; Indels 1; Gaps 1;  
  
Qy 1 AACAGGCTCTCTCTCAGTGTATCAACTTTTGACACTTGTGCGATCGGTGATGCTGTCT 60  
Db 3 AACAGGCTCTCTCTCAGTGTATCAACTTTTGACACTTGTGCGATCTCTGTGCTGTCT 62  
Qy 61 GCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTTGGCCGCGCAGCTGCTGTCTCAT 120  
Db 63 GCAGAAATCTATGAGTTTTTCCCTTATGGGAGCTTTTGGCCGCGCAGCTGCTGTCTCAT 122  
Qy 121 TGCCCTGTGGGCCAGGAGCAAAATGCGTGTGCCCATCAACACCCGGTGCAGCTTGAGGT 180  
Db 123 TGCCCTGTGGGCCAGGAGCAAAATGCGTGTGCCCATCAACACCCGGTGCAGCTTGAGGT 182  
Qy 181 GTCCAACTTCAGAGCCGTACATCGTCAACCCGACCTTTATGCTGCGCAGAGGCGCAG 240  
Db 183 GTCCAACTTCAGAGCCGTACATCGTCAACCCGACCTTTATGCTGCGCAGAGGCGCAG 242  
Qy 241 CCTTGACAGATAACAAACACAGACGTCGGGCTCATCGGGGAGAAACTGTTCGAGAGTCAG 300  
Db 243 CCTTGACAGATAACAAACACAGACGTCGGGCTCATCGGGGAGAAACTGTTCGAGAGTCAG 302  
Qy 301 TGCTAAGGATCAGTGTCTACCTGTATGAACAGAGTGTCTCAACTTCACTTCAAGACGTTCT 360  
Db 303 TGCTAAGGATCAGTGTCTACCTGTATGAACAGAGTGTCTCAACTTCACTTCAAGACGTTCT 362  
Qy 361 GCTCCCCCAGTCAGACAGGTTCCGGCCCTACATCGCAGAGGTGTGCTTCTCTGACCAA 420  
Db 363 GCTCCCCCAGTCAGACAGGTTCCAGCCCTACATGCGAGGTTGGTACCTTTCTCTGACCAA 422  
Qy 421 ACTCAGCAATCAGCTCAGCTCCTGTCTACATCAGTGTGACGACAGCAACATCCAGAGAA 480  
Db 423 ACTCAGCAATCAGCTCAGCTCCTGTCTACATCAGGCGTGAACAGCAACATCCAGAGAA 482  
Qy 481 TGTGAGAAAGCTGAAGGAGACAGTGAAAGAACTTTGGAGAGCGGAGAGATCAAGCGAT 540  
Db 483 TGTGAGAAAGCTGAAGGAGACAGTGAAAGAACTTTGGAGAGCGGAGAGATCAAGCGAT 542  
Qy 541 CGGGGAATCGAATCTGCTGTTTATGTCTCTGAGAAATGCTTGCCTGTGAGCGAGAGAG 600  
Db 543 TGGGGAATCTGGACCTGCTGTTTATGTCTCTGAGAAATGCTTGCCTGTGAGCGAGAGAG 602  
Qy 601 CTAGAAACGAGAACTGCTCCTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 660  
Db 603 CTAGAAACGAGAACTGCTCCTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 662  
Qy 661 GACTTTTTTACTAAGGAAAGTGAAGAGCTAACTGCTCAACATCAATTAGAGATTTTCACAT 720  
Db 663 GACTTTTTTACTAAGGAAAGTGAAGAGCTAACTGCTCAACATCAATTAGAGATTTTCACAT 722  
Qy 721 GAAACCTGGCTCAGTGTGAAAGAGAAATAGTGTCAAGTTGTCCATGAGACCAAGAGGTAGA 780  
Db 723 GAAACCTGGCTCAGTGTGAAAGAGAAATAGTGTCAAGTTGTCCATGAGACCAAGAGGTAGA 782
```

Qy	781	CTTGATTAACCAACAAGATTCAATGACAATAATTTTATGTCAATGATAAATGCACAGAATAA	840
Db	783	CTTGATAAACCAACAAGATTCAATGACAATAATTTTATGTCACTGATGATACAACAAGAATAA	842
Qy	841	AGTATGTACTTTAAAAAATTTGTTGAAGAGAGGTTACCTCTCATTCCTCTAGAAGAAAAG	900
Db	843	ATAATGTACTTTAAAAAATTTGTTGAAGAGAGGTTACCTCTCATTCCTTTAGAAAAAATAAG	902
Qy	901	CCATGTGTAACTTCATTTCCATAACCATACTTTATATATGATGAAGTTTTATTATATATAAGT	960
Db	903	CTTATGTGAACCTTCATTTCCATATCCAATATTTTATATATGATGAAGTTTTATTATATAAGT	962
Qy	961	ATACATTTTATTTATGTGCAGTTTATTAATATGGATTTTATTTATAGAAAAATTAATCTGATG	1020
Db	963	ATACATTTTATTTATGTGCAGTTTATTAATATGATTTTATTTATAGAAAAATTAATCTGCTA	1022
Qy	1021	TTCGATATTTTCAGTATAAAGCAAATAATTTATTTGATAATAACTATAGAAAACAAGATATCT	1080
Db	1023	TTTGATATTT-AGTATAGGCCAATAATATTTTATGACAAATAACTATGGAACAAGATATCT	1081
Qy	1081	TAGCCTTTAATAAACACATGAATATCATAAA	1111
Db	1082	TAGCCTTTAATAAACACATGGATATCATAAA	1112

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RESULT 3
US-11-177-987-42
; Sequence 42, Application US/11177987
; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors and Uses Thereof
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 42
; LENGTH: 5935
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-11-177-987-42

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Qy	748	TAGTGTCAAGTTGTCCTCAGTAGAGCCAGAGGTAGAGCTTTGATAA CCAAAAGATTCAATTGACA	807
Db	5461	TAGTGTCAAGTTGTCCTCAGTAGAGCCAGAGGTAGAGCTTTGATAA CCAAAAGATTCAATTGACA	5520
Qy	808	ATATTTTATTGTTCATTGTAGATAATCCAAACAGAAAAAGATGTAC TTTTAAAAAATTTGTTTGA	867
Db	5521	ATATTTTATTGTTCATTGTAGATAATCCAAACAGAAAAAGATGTAC TTTTAAAAAATTTGTTTGA	5580
Qy	868	AGGAGGTTACCTCTCTCAATTCCTCTAGAAAGAAAGCCCTATGTAA CTTCAATTTCCATAACCAA	927
Db	5581	AGGAGGTTACCTCTCTCAATTCCTCTAGAAAGAAAGCCCTATGTAA CTTCAATTTCCATAACCAA	5640
Qy	928	TACTTTTATATATCTGAAGTTTATTATTATTATAAGATATACATT TTTTATTTATGTGCAGTTTATTA	987
Db	5641	TACTTTTATATATCTGAAGTTTATTATTATTATAAGATATACATT TTTTATTTATGTGCAGTTTATTA	5700
Qy	988	ATATGGATTTTATTATAGAAAAATTA TCTGATGTTTGATATTTGAGTATATAAGCAAAATAAT	1047
Db	5701	ATATGGATTTTATTATAGAAAAATTA TCTGATGTTTGATATTTGAGTATATAAGCAAAATAAT	5760
Qy	1048	ATTATTGATATAATTAACATATAGAAA CAAGATATCTTAGGCTTTTAATAAACACATGAATATCA	1107
Db	5761	ATTATTGATATAATTAACATATAGAAA CAAGATATCTTAGGCTTTTAATAAACACATGAATATCA	5820
Qy	1108	TAAA 1111	
Db	5821	TAAA 5824	

RESULT 4

US-11-177-987-8

; Sequence 8, Application US/11177987

; Publication No. US20050271619A1

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Renauld, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Induc

; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5664

; CURRENT FILING DATE: 2005-07-07

; PRIOR APPLICATION NUMBER: US/11/177,987

; PRIOR FILING DATE: 2005-07-07

; PRIOR APPLICATION NUMBER: US/09/626,617

; PRIOR FILING DATE: 2000-07-27

; PRIOR APPLICATION NUMBER: US09/419,568

; PRIOR FILING DATE: 1999-10-18

; PRIOR APPLICATION NUMBER: US09/354,243

; PRIOR FILING DATE: 1999-07-16

; PRIOR APPLICATION NUMBER: US09/178,973

; PRIOR FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 43

; SEQ ID NO 8

; LENGTH: 7445

; TYPE: DNA

; ORGANISM: Mus musculus

; FEATURE:

US-11-177-987-8

Query Match	50.0%	Score 555.2	DB 7	Length 7445
Best Local Similarity	96.0%	Pred. No. 3.1e-120		
Matches 580; Conservative	0	Mismatches 23	Indels 1	Gaps 1
Qy	508	AAAGCTTGGAGAGCGGAGAGATCAAAGCGATCGGGGAACTCGACTCGTGTGTTTATGTC	567	
Db	6535	ATAGCTTGGAGAGATGGAGAGATCAAGCGATTGGGGAACCTGGACTGCTGTTTATGTC	6594	
Qy	568	TCTGAGAAATGCTTTCGCTCTGACGCGAGAACTAGAAAAAGAAAGCTGCTTCCTTCCT	627	
Db	6595	TCTGAGAAATGCTTTCGCTCTGACGCGAGAACTAGAAAAAGAAAGCTGCTTCCTTCCT	6654	
Qy	628	GCCTTCTAAAAAGAAACAATTAAGATTCCTGAATCGACTTTTTTACTAAGAAAGTCGAGAA	687	
Db	6655	GCCTTCTAAAAAGAAACAATTAAGATTCCTGAATCGACTTTTTTACTAAGAAAGTCGAGAA	6714	



US-11-177-987-25  
; Sequence 25, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 25  
; LENGTH: 690  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-11-177-987-25  
Query Match 36.7%; Score 407.6; DB 7; Length 690;  
Best Local Similarity 76.0%; Pred. No. 3.5e-86;  
Matches 503; Conservative 0; Mismatches 159; Indels 0; Gaps 0;  
QY 7 CTCCTCTCAGTATCAACTTTTGACACTTGTGCGATCGGTGATGGCTGTCTCTGCAGAA 66  
DB 29 CTCCTTCCCAGTCACAGTTGCTCGAGTTAGAAATGCTGCAATGCGCCCTGCAGAA 88  
QY 67 ATCTATGAGTTTTCCCTTATGGGACTTTGGCGGCAGCTGCTGCTTCTCATTTGCCCT 126  
DB 89 ATCTGTGAGCTTCTTCTTATGGGACCTTGGCCACAGCTGCTCTTCTTGGCCCT 148  
QY 127 GTGGGCCAGGAGCAATCGCTGCCCATCAACCCGGTGAAGCTTCAGGTGTCCAA 186  
DB 149 CTTGGTACAGGAGGACAGCTGGCGCCCATCAGTCCCATCGAGGCTTGACAAAGTCCAA 208  
QY 187 CTTCCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGGAGCGACCTTGC 246  
DB 209 CTTCCAGCAGCGCTATATACCAACCGCACCTTTCATGCTGCTTAAGGAGCTAGCTTGC 268  
QY 247 AGATAACAAACAGACGCTCGGCTCATCGGGGAGAACTGTTCCGAGGAGTCAAGTGTAA 306  
DB 269 TGATAACAAACAGACGCTTCTCTCATTTGGGGAGAACTGTTCCACGAGTCAAGTATGAG 328  
QY 307 GGATCAGTGTACTGTATGAGCAGGTGCTCACTTCCACCTCGAGACATTTCTGCTCC 366  
DB 329 TGAGCGCTGTCTATCTGATGAGCAGGTGCTCACTTCCACCTTGAAGAGTGTGTTCC 388  
QY 367 CCAGTCAGACAGGTTCCGGGCTTACATGACGAGGTTGGTGCCTTTCTCTGACCAAACTCAG 426  
DB 389 TCAATCTGATAGTTTCAGGCTTATATGACGAGGTTGGTGCCTTCTTGGCCAGGCTCAG 448  
QY 427 CAATCAGCTCAGTCTGTTCATATCAGTGTGACGACCAAACTCCAGAAAGATGTTCAG 486  
DB 449 CAACAGGCTAAGCACATGTCTATATGAAGGTGATGACCTGCATATCCAGAGGAATGTCA 508  
QY 487 AAGCTCAAGGACAGTGAAGAGCTTGGAGAGCGGAGAGATCAAGCGATCCGGGA 546  
DB 509 AAGCTCAAGGACAGTGAAGAGCTTGGAGAGGTTGGAGAGATCAAGCAATTTGAGA 568  
QY 547 ACTGGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTGTGACGAGAGAAAGCTAGAA 606  
DB 569 ACTGGATTGCTGTTTATGCTCTGAGAAATGCTGCTGATTTGACGAGCAAGCTGAAA 628  
QY 607 AACGAGAACTGCTCTTCTCTGCTTCTTAAAGAAACAAATAAGATCCCTGAATGGACTTTT 666

DB 629 AATGAATAACTAACCCCTTTCCTCTAGAAATAACAATTAGATGCCCCAAAGCGATTT 688  
QY 667 TT 668  
DB 689 TT 690  
RESULT 7  
US-11-177-987-18  
; Sequence 18, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 18  
; LENGTH: 418  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-11-177-987-18  
Query Match 19.5%; Score 217.2; DB 7; Length 418;  
Best Local Similarity 73.2%; Pred. No. 9.9e-42;  
Matches 293; Conservative 0; Mismatches 103; Indels 4; Gaps 1;  
QY 352 AGACATTCCTGCTCCCCCAGTCACACAGGTTTCGGGCCCTACATCAGGAGTGGTGCCTTT 411  
DB 1 AGAAGTGTGTTCCCTCAATCTGATAGTTTCAGGCTTATATCAGGAGTGGTGCCTTT 60  
QY 412 CTTGACCAAACTCAGCAATCAGCTCAGCTCCTGTCTCATCAGTGGTGAACGACGAGAACAT 471  
DB 61 CTTGGCCAGGCTCAGCAACAGGCTAAGCACATGTCATATTGAAGGTGATGACCTGCATAT 120  
QY 472 CAGAAAGATGTGAGAAGGCTGAAGGAGACAGTGAAGAAAGCTTGGAGAGCGGAGAGAT 531  
DB 121 CCAGAGGAATGTGCAAAAGCTGAAGGACACAGTGAAGAAAGCTTGGAGAGTGGAGAGAT 180  
QY 532 CAAAGCGATCGGGGAATCGACCTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGC 591  
DB 181 CAAAGCAATTTGAGAAAGTGGATTTGCTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGC 240  
QY 592 GAGAAAGCTAGAAACGAAAGCTGCTCTTCTGCTGCTTCTTAAAGAAACAAATAGAT 651  
DB 241 AGAGCAAGCTGAAGAAATGAATAACTAACCCCTTTCCTGCTAGAAATAACAATTAGAT 300  
QY 652 CCTGTAATGAGCTTTTTTTT-----ACTAAAGAAAGTGAAGAGCTAACGTCACCATCATTA 707  
DB 301 GCCCAAAGCGATTTTTTTTTTAAACCAAAAGGAAGATGGGAAGCCAAATCTCATATGATGG 360  
QY 708 GAAGATTTTCATGAACCTGGCTCAGTTGAAGAGAGAAA 747  
DB 361 GTGGATTCCAATGAACCCCTCGCTGTTAGTTACAAAGAAA 400  
RESULT 8  
US-11-177-987-26  
; Sequence 26, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:





; Sequence 105, Application US/11121086  
; Publication No. US20050266459A1  
; GENERAL INFORMATION:  
; APPLICANT: POULSEN, TIM S.  
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES  
; FILE REFERENCE: 09138.6000-00000  
; CURRENT APPLICATION NUMBER: US/11/121,086  
; PRIOR FILING DATE: 2005-05-04  
; PRIOR FILING DATE: 2005-05-04  
; PRIOR FILING DATE: 2004-05-04  
; NUMBER OF SEQ ID NOS: 107  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 105  
; LENGTH: 171486  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-11-121-086-105

Query Match 5.0%; Score 55.6; DB 7; Length 171486;  
Best Local Similarity 54.8%; Pred. No. 0.012;  
Matches 172; Conservative 0; Mismatches 139; Indels 3; Gaps 3;  
QY 799 TCATTGACATATTTTATGTCCTGATGTAATGCAACAGAAAAGTATGTAATTTTAA 858  
Db 16418 TAATATATATATTTATTAATTTATATATATATATATATATATATATATATAT 16477  
QY 859 TTGTTTGAAGGAGGTTCCTGCTCATTCTCTAGAGAAAAGCTGATGTAATTTTC 918  
Db 16478 TATATATATATTTATTTACATATTTATATATATATATATATATATATATAT 16536  
QY 919 CATAACCAATFACCTTTATATATGTAAGTTTATTTATTTATATA-AGTATACATTTTATG 977  
Db 16537 AATATATATACAT 16596  
QY 978 CAGTTTATTAATGAGTTTATTTATAGAAAATTAATCTGATGTTGATTTGAGTATA 1037  
Db 16597 TTATATATTTAT 16656  
QY 1038 AGCAATAATTTATGATATATATATATATATATATATATATATATATATATAT 1097  
Db 16657 TATTTACATATTAAT-ATATATATATATATATATATATATATATATATATAT 16715  
QY 1098 ATGAATATCATATA 1111  
Db 16716 TATTATATATAA 16729

RESULT 12  
US-10-995-561-13473  
; Sequence 13473, Application US/10995561  
; Publication No. US2005027054A1  
; GENERAL INFORMATION:  
; APPLICANT: CARGILL, Michele et al.  
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH  
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF  
; FILE REFERENCE: CL001559  
; CURRENT APPLICATION NUMBER: US/10/995,561  
; CURRENT FILING DATE: 2004-11-24  
; NUMBER OF SEQ ID NOS: 85702  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 13473  
; LENGTH: 394468  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: misc feature  
; LOCATION: (1)- (394468)  
; OTHER INFORMATION: n = A,T,C or G, or insertion/deletion polymorphism (see Tables 1-  
US-10-995-561-13473

Query Match 5.0%; Score 55.6; DB 6; Length 394468;

Best Local Similarity 50.0%; Pred. No. 0.018;  
Matches 168; Conservative 1; Mismatches 160; Indels 7; Gaps 1;  
QY 782 TTGATAACCAACAAAGATTTCATTGACAATATTTTATTGTTCATTGATTAATGCAACAGAAAA 841  
Db 340523 TTATATAATATAAATATAAATATATATTTGTTTATATATATATTTTATAAACAATAATATG 340582  
QY 842 GTATGTACTTTAAAAAATTTGTTGAAGGAGGTACCTCTCAATTCCTCTAGA-----A 894  
Db 340583 TTTTATACATATAAAAAATTTGTATATAAAATTTATATCTATATGAAATATAGATATATTT 340642  
QY 895 GAAAGCCTATGTAACCTTCATTTCACCAACCAATCTTTATATATCTAGTTTATTTATTT 954  
Db 340643 ATAAATATATATATATATTTATGTATATAATATATGCAATTTATATATTTATGTATAAATATA 340702  
QY 955 ATAAGTATACATTTTATTTATGTCAGTTTATTAATATGGAATTTATTTATAGAAAAATTTAT 1014  
Db 340703 TGCATTTATATATTTATGTATAAATATATATGCAATTTATTTATGTATAAATATATGCA 340762  
QY 1015 CTGATGTTGATATTTGAGTATATAAGCAATAATATTTATGTATATAAATCACTATAGAAACAAG 1074  
Db 340763 TTTATATATTTATGTATAAATATATGCAATTTATATATATTTATGTATAAATATATATGCA 340822  
QY 1075 ATATCTTAGGCTTTAATAACACATGCAATATCATAA 1110  
Db 340823 TATATTTATGTATAAATATATATGCAATTTATATATTTA 340858

RESULT 13  
US-10-995-561-13313/c  
; Sequence 13313, Application US/10995561  
; Publication No. US2005027054A1  
; GENERAL INFORMATION:  
; APPLICANT: CARGILL, Michele et al.  
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH  
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF  
; FILE REFERENCE: CL001559  
; CURRENT APPLICATION NUMBER: US/10/995,561  
; CURRENT FILING DATE: 2004-11-24  
; NUMBER OF SEQ ID NOS: 85702  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 13313  
; LENGTH: 26772  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-10-995-561-13313

Query Match 5.0%; Score 55.2; DB 6; Length 26772;  
Best Local Similarity 50.7%; Pred. No. 0.0061;  
Matches 152; Conservative 2; Mismatches 145; Indels 1; Gaps 1;  
QY 767 AGACCAGAGGTAGACTTTGATTAACCAACAGATTTCATTGACAATTTTATTGTCATTGAT 826  
Db 21931 ATATAAATRTATTTATTTATATATATATATATATATTTCTATATAAATATATATATATAT 21872  
QY 827 AATGCAACAGAAAAAGTATGTAATTTAAAAAATTTGTTGAAAGGAGGTACCTCTCATTC 886  
Db 21871 ATAAATTTCTATATAAATATATATGTTATTTATTTATATAGATACATATATAK-TATATA 21813  
QY 887 CTCTAGAGAAAGCCTATGTAACCTTCATTTCATCAACCAATCTTTATATATATGTAAGTT 946  
Db 21812 TAATATTTATTTATTTATATATATATTAATTAATTAATTAATTAATTAATTAATTAATAA 21753  
QY 947 TATTTATTAAGTATACATTTTATTTATGTCAGTTTATTAATTAATGATGATTTATATAGA 1006  
Db 21752 TTATATATAAATATATAATTTATTTATTTATTTATTTATTTATTTATTTATTTATTTAT 21693  
QY 1007 AAAATTTCTGATGTTGATTTTGGATATAAAGCAAAATTAATTTTATGTAATAAATCTATA 1066  
Db 21692 TATAATATAAATTAATTTATTTATTTATTTATTTATTTATTTATTTATTTATTTATTTAT 21633



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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 48.1106 Seconds  
(without alignments)  
7442.822 Million cell updates/sec

Title: US-09-751-797-24  
Perfect score: 690  
Sequence: 1 tgcacagcagaattctcag.....gatgcccacagcgattttt 690

Scoring table: IDENTITY NUC  
Gapop 10.0 , Gapext 1.0

Searched: 4168288 seqs, 259477437 residues

Total number of hits satisfying chosen parameters: 8336576

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Published Applications\_NA\_New.\*  
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2: /cgn2\_6/ptodata/2/pubpna/US06\_NEW\_PUB.seq.\*  
3: /cgn2\_6/ptodata/2/pubpna/US07\_NEW\_PUB.seq.\*  
4: /cgn2\_6/ptodata/2/pubpna/PCT\_NEW\_PUB.seq.\*  
5: /cgn2\_6/ptodata/2/pubpna/US09\_NEW\_PUB.seq.\*  
6: /cgn2\_6/ptodata/2/pubpna/US10\_NEW\_PUB.seq.\*  
7: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq.\*  
8: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq2.\*  
9: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq3.\*  
10: /cgn2\_6/ptodata/2/pubpna/US60\_NEW\_PUB.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	690	100.0	690	7	US-11-177-987-25
2	676	98.0	1152	7	US-11-102-240-153
3	409.2	59.3	1119	7	US-11-177-987-7
4	407.6	59.1	1111	7	US-11-177-987-9
5	317	45.9	418	7	US-11-177-987-18
6	258	37.4	4797	7	US-11-177-987-26
7	127.6	18.5	5935	7	US-11-177-987-42
8	126	18.3	7445	7	US-11-177-987-8
9	38.8	5.6	1816	6	US-10-750-185-40262
10	36.4	5.3	152335	7	US-11-121-086-73
11	36	5.2	3485	6	US-10-821-234-300
C 12	34.4	5.0	1437	6	US-10-750-185-43510
C 13	33.8	4.9	3044	6	US-10-750-185-31621
14	32.6	4.7	2673	6	US-10-750-185-43708
15	32.2	4.7	2126	6	US-10-750-185-50850
C 16	32	4.6	3641	7	US-11-113-424-19
C 17	32	4.6	3692	7	US-11-113-424-21
C 18	31.6	4.6	86081	6	US-10-995-561-13246
C 19	31.4	4.6	2697	6	US-10-750-185-59603
C 20	31	4.5	600	6	US-10-750-185-1430
C 21	31	4.5	873	6	US-10-750-185-60900
C 22	31	4.5	1183	6	US-10-995-561-504
23	31	4.5	2138	6	US-10-995-561-500

ALIGNMENTS

RESULT 1

US-11-177-987-25  
; Sequence 25, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors and Uses Thereof  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; PRIOR FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 25  
; LENGTH: 690  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-11-177-987-25  
Query Match 100.0%; Score 690; DB 7; Length 690;  
Best Local Similarity 100.0%; Pred. No. 3.2e-215;  
Matches 690; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 1 TGCACAGCAGAAATCTTTCAGAACAGAGTTCTCTCCCACTCACCAGTTGCTCGAGTTAG 60  
Db 1 TGCACAGCAGAAATCTTTCAGAACAGAGTTCTCTCCCACTCACCAGTTGCTCGAGTTAG 60  
Qy 61 AATTGCTGCAATGGCGCCCTCGAGAAATCTGTAGCTCTTCTTATGGGACCTGG 120  
Db 61 AATTGCTGCAATGGCGCCCTCGAGAAATCTGTAGCTCTTCTTATGGGACCTGG 120  
Qy 121 CCACCACTGCTCCCTCTTCTTGGCCCTCTTGTGTACAGGAGGAGCAGCTCGCCCATCA 180  
Db 121 CCACCACTGCTCCCTCTTCTTGGCCCTCTTGTGTACAGGAGGAGCAGCTCGCCCATCA 180  
Qy 181 GTCCCACTGAGGCTTGCAAGTCCAACTTTCAGAGGCCCTTATATCACCAGCGACT 240  
Db 181 GTCCCACTGAGGCTTGCAAGTCCAACTTTCAGAGGCCCTTATATCACCAGCGACT 240

Db 181 GCTCCACTGCAGGCTTGACAAGTCCAACTCCAGCAGCCCTATATCACCAACCGCACCT 240  
Qy 241 TCATGCTGGCTAAGGAGGCTAGCTTGGCTGATATAACAACAGAGCTTCGTCATTTGGGG 300  
Db 241 TCATGCTGGCTAAGGAGGCTAGCTTGGCTGATATAACAACAGAGCTTCGTCATTTGGGG 300  
Qy 301 AGAACTGTTCCACGGAGTCAAGTATGATGAGTGAAGCGCTGCTATCTGATGAAGCAGGTGCTGA 360  
Db 301 AGAACTGTTCCACGGAGTCAAGTATGATGAGTGAAGCGCTGCTATCTGATGAAGCAGGTGCTGA 360  
Qy 361 ACTTCACCCCTTGAAGAAGTGTGTTCCCTCAATCTGATAGGTTCCAGCCTTATATGCAAG 420  
Db 361 ACTTCACCCCTTGAAGAAGTGTGTTCCCTCAATCTGATAGGTTCCAGCCTTATATGCAAG 420  
Qy 421 AGGTGCTGCTTCCCTGCGCCAGGCTCAGCAACAGGCTAAGCAATGTCATATTAAGAGGTG 480  
Db 421 AGGTGCTGCTTCCCTGCGCCAGGCTCAGCAACAGGCTAAGCAATGTCATATTAAGAGGTG 480  
Qy 481 ATGACCTGCATATCCAGAGGATGTCGAAAGCTGAAGGACACAGTGAAGAAGCTTGGAG 540  
Db 481 ATGACCTGCATATCCAGAGGATGTCGAAAGCTGAAGGACACAGTGAAGAAGCTTGGAG 540  
Qy 541 AGAGTGGAGAGATCAAAAGCAATGGAGAACTGGATTTGCTTTATGTTCTCTGAGAAATG 600  
Db 541 AGAGTGGAGAGATCAAAAGCAATGGAGAACTGGATTTGCTTTATGTTCTCTGAGAAATG 600  
Qy 601 CCTGATTTGACAGAGCAAAAGCTGAAATAATGAATACTAACCCCTTCCCTGCTAGAA 660  
Db 601 CCTGATTTGACAGAGCAAAAGCTGAAATAATGAATACTAACCCCTTCCCTGCTAGAA 660  
Qy 661 ATACAAATAGATGCCCCAAGCGATTTTT 690  
Db 661 ATACAAATAGATGCCCCAAGCGATTTTT 690

RESULT 2

US-11-102-240-153  
; Sequence 153, Application US/11102240  
; Publication No. US20050260647A1  
; GENERAL INFORMATION:  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Godowski, Paul J.  
; APPLICANT: Grimaldi, Christopher J.  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Wood, William I.  
; TITLE OF INVENTION: ANTIBODIES TO POLYPEPTIDES ENCODED BY A NUCLEIC ACID UNDEREXPRESS  
; TITLE OF INVENTION: ESOPHAGEAL TUMOR  
; FILE REFERENCE: P3230R1C106C  
; CURRENT APPLICATION NUMBER: US/11/102,240  
; CURRENT FILING DATE: 2005-04-08  
; PRIOR APPLICATION NUMBER: 10/063662  
; PRIOR FILING DATE: 2002-05-07  
; PRIOR APPLICATION NUMBER: 10/006867  
; PRIOR FILING DATE: 2001-12-06  
; PRIOR APPLICATION NUMBER: PCT/US00/23328  
; PRIOR FILING DATE: 2000-08-24  
; PRIOR APPLICATION NUMBER: 60/170262  
; PRIOR FILING DATE: 199-12-09  
; NUMBER OF SEQ ID NOS: 170  
; SEQ ID NO 153  
; LENGTH: 1152  
; TYPE: DNA  
; ORGANISM: Homo Sapien  
US-11-102-240-153

Query Match 98.0%; Score 676; DB 7; Length 1152;  
Best Local Similarity 100.0%; Pred. No. 1.6e-210; Indels 0; Gaps 0;  
Matches 676; Conservative 0; Mismatches 0

Qy 15 CTTCAAGACAGGTTCTCCCTCCCGAGTCACCGAGTTCGAGTTGAGAAATGTTCTGCAATG 74  
Db 1 CTTCAAGACAGGTTCTCCCTCCCGAGTCACCGAGTTCGAGTTGAGAAATGTTCTGCAATG 60

Qy 75 GCGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGGCCACCAGAGCTGCCTC 134  
Db 61 GCGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGGCCACCAGAGCTGCCTC 120  
Qy 135 CTTCTCTTGGCCCTCTTTGGTACAGGGAGGAGCGAGCTGCGCCCATCAGCTCCCATGCGAGG 194  
Db 121 CTTCTCTTGGCCCTCTTTGGTACAGGGAGGAGCGAGCTGCGCCCATCAGCTCCCATGCGAGG 180  
Qy 195 CTTGACAAAGTCCAACTTCCAGAGCCCTATATCACCAACCGCACCTTCATGCTGGCTAAG 254  
Db 181 CTTGACAAAGTCCAACTTCCAGAGCCCTATATCACCAACCGCACCTTCATGCTGGCTAAG 240  
Qy 255 GAGCTAGCTTGGCTGATAAACAACAGAGCTTCCTCATTTGGGGAGAAACTGTTCCAC 314  
Db 241 GAGCTAGCTTGGCTGATAAACAACAGAGCTTCCTCATTTGGGGAGAAACTGTTCCAC 300  
Qy 315 GAGTCAAGTATGAGTGAGCGCTGCTATCTGATGAAGCAGGCTGCTCAACCCCTTGAA 374  
Db 301 GAGTCAAGTATGAGTGAGCGCTGCTATCTGATGAAGCAGGCTGCTCAACCCCTTGAA 360  
Qy 375 GAAGTGTGTTCCCTCAATCTGATAGGTTCCAGCCTTATATGCAAGAGGTGGTCCCTTC 434  
Db 361 GAAGTGTGTTCCCTCAATCTGATAGGTTCCAGCCTTATATGCAAGAGGTGGTCCCTTC 420  
Qy 435 CTGSCCAGGCTCAGCAACAGGCTAAGCACATGTCAATTTGAAGGTGATGACCTGCATATC 494  
Db 421 CTGSCCAGGCTCAGCAACAGGCTAAGCACATGTCAATTTGAAGGTGATGACCTGCATATC 480  
Qy 495 CAGAGGAATGTGCAAAAGCTGAAAGGACACAGTGAAGAAAGCTTTGGAGAGGTGGAGAGATC 554  
Db 481 CAGAGGAATGTGCAAAAGCTGAAAGGACACAGTGAAGAAAGCTTTGGAGAGGTGGAGAGATC 540  
Qy 555 AAAGCAATGGAGAACTGGATTTGCTGTTTATGCTCTGAGAAATGCTGCAATTTGACCA 614  
Db 541 AAAGCAATGGAGAACTGGATTTGCTGTTTATGCTCTGAGAAATGCTGCAATTTGACCA 600  
Qy 615 GAGCAAGCTGAAATAATGAATACTAACCCCTTCCCTGCTAGAAAATAACAATTAGATG 674  
Db 601 GAGCAAGCTGAAATAATGAATACTAACCCCTTCCCTGCTAGAAAATAACAATTAGATG 660  
Qy 675 CCCCCAAGCGATTTTT 690  
Db 661 CCCCCAAGCGATTTTT 676

RESULT 3

US-11-177-987-7  
; Sequence 7, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-11-177-987-7



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; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 18
; LENGTH: 418
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-11-177-987-18

Query Match      45.9%; Score 317; DB 7; Length 418;
Best Local Similarity 100.0%; Pred. No. 1.6e-93;
Matches 317; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 374 AGAAGTGTCTCCCTCAATCTGATAGTTCCAGCCTTATATGCGAGGTTGGTCCCTT 433
Db 1 AGAAGTGTCTCCCTCAATCTGATAGTTCCAGCCTTATATGCGAGGTTGGTCCCTT 60

Qy 434 CCTGCCAGGCTCAGCAACAGCTAAGCACATGTATATGAAGTGATGACCTGCGATAT 493
Db 61 CCTGCCAGGCTCAGCAACAGCTAAGCACATGTATATGAAGTGATGACCTGCGATAT 120

Qy 494 CCAGAGGAATGTCAAAAGCTGAAGGACACAGTGAAGGCTGAGAGAGTGGAGAGAT 553
Db 121 CCAGAGGAATGTCAAAAGCTGAAGGACACAGTGAAGGCTGAGAGAGTGGAGAGAT 180

Qy 554 CAAAGCAATGGAGACTGGATTTGCTGTTATGTCTCTGAGAAATGCGTGCATTTGACC 613
Db 181 CAAAGCAATGGAGACTGGATTTGCTGTTATGTCTCTGAGAAATGCGTGCATTTGACC 240

Qy 614 AGAGCAAGCTGAAAAATGAATACTAACCCCTTTCCCTGCTAGAAATAACAATTAGAT 673
Db 241 AGAGCAAGCTGAAAAATGAATACTAACCCCTTTCCCTGCTAGAAATAACAATTAGAT 300

Qy 674 GCCCCAAAGCGATTTT 690
Db 301 GCCCCAAAGCGATTTT 317

RESULT 6
US-11-177-987-26
; Sequence 26, Application US/11/177987
; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renault, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 26
; LENGTH: 4797
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-11-177-987-26
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Query Match      37.4%; Score 258; DB 7; Length 4797;
Best Local Similarity 100.0%; Pred. No. 1.2e-73;
Matches 258; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TGCACAAGCAGAAATCTTTCAGAACAGAGTTCTCTCCCTCCAGTCACCAAGTTGCTCGAGTTAG 60
Db 1 TGCACAAGCAGAAATCTTTCAGAACAGAGTTCTCTCCCTCCAGTCACCAAGTTGCTCGAGTTAG 60

Qy 61 AATTGTCTGCAANTGGCCGCTCGAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGG 120
Db 61 AATTGTCTGCAANTGGCCGCTCGAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGG 120

Qy 121 CCACCAAGTCCCTCTCTTCTTGGCCCTCTTGTGTACAGGAGGAGGAGTCTGCGCCCATCA 180
Db 121 CCACCAAGTCCCTCTCTTCTTGGCCCTCTTGTGTACAGGAGGAGGAGTCTGCGCCCATCA 180

Qy 181 GCTCCCACTGCGAGCTTTGACAAAGTCCAACCTTCCAGCAGCCCTATATCACCACCGCACCT 240
Db 181 GCTCCCACTGCGAGCTTTGACAAAGTCCAACCTTCCAGCAGCCCTATATCACCACCGCACCT 240

Qy 241 TCATGCTGGCTTAAGGAGG 258
Db 241 TCATGCTGGCTTAAGGAGG 258

RESULT 7
US-11-177-987-42
; Sequence 42, Application US/11/177987
; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renault, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 42
; LENGTH: 5935
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-11-177-987-42

Query Match      18.5%; Score 127.6; DB 7; Length 5935;
Best Local Similarity 72.2%; Pred. No. 5.1e-31;
Matches 166; Conservative 0; Mismatches 64; Indels 0; Gaps 0;

Qy 29 CTCCTTCCCAGTCACCAAGTTGCTCGAGTTAGAATTGTCTGCAATGGCCGCTCGAGAA 88
Db 356 CTCCTTCCCAGTCACCAAGTTGCTCGAGTTAGAATTGTCTGCAATGGCCGCTCGAGAA 415

Qy 89 ATCTGTGAGCTCTTTTCTTATGGGACCCCTGCGCCACAGCTGCGCTCTCTTCTTGGCCCT 148
Db 416 ATCTGTGAGTTTTCCTTATGGGACCTTGGCCGCGCCAGCTGCTGCTTCTCATTTGCCCT 475

Qy 149 CTTGGTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCATCTGCAAGGTTGACAAGTCAA 208
Db 476 GTGGGCCCGAGGAGCAAAATGCGTCCCATCAACACCGGTCGAAGCTTGAGGTGTCCAA 535

Qy 209 CTTCCAGCAGCCCTATATCACCACCGCACCTTCTATGCTGCTAAGGAGG 258
Db 536 CTTCCAGCAGCCCTATATCAGTCAACCGCACCTTTATGCTGCGCCAGGAGG 585
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; CURRENT APPLICATION NUMBER: US/10/821,234
; CURRENT FILING DATE: 2004-04-07
; PRIOR APPLICATION NUMBER: US 60/462,047
; PRIOR FILING DATE: 2003-04-07
; NUMBER OF SEQ ID NOS: 1704
; SOFTWARE: pt_SEQ_genes Version 1.0
; SEQ ID NO 300
; LENGTH: 3485
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-821-234-300

Query Match          5.2%; Score 36; DB 6; Length 3485;
Best Local Similarity 50.6%; Pred. No. 0.31;
Matches 87; Conservative 0; Mismatches 85; Indels 0; Gaps 0;

Qy 343 TGATGAAGCAGGTGCTGAACTTCACTTGAAGAGTGTGCTTCCCTCAATCTGATAGGT 402
Db 1903 TGGTGCAGGAGGAGAGCCCTTCAGCGAGGAAGCTACACTTTTCCACCAAGAACTGGTGC 1962

Qy 403 TCGAGCCTTATGACGAGAGTGTGCTTCCCTTCTGGCCAGGCTCAGCAACAGGCTAAGCA 462
Db 1963 TGCAGCGAGAGGTGGAGGTGGAGATGGAACAGCCGCGCAACTTTTATCGGCT 2022

Qy 463 CATGTCTATTGAAGGTGATGACCTGCATATCCAGAGGAATGTGCAAAAGCT 514
Db 2023 GGCTGCATCGACGGTGCCAACTGTCCGTCTCTGCTGGTGAGACGCGCT 2074

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RESULT 12
US-10-750-185-43510/c
; Sequence 43510, Application US/10750185
; Publication No. US20050260603A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-2
; CURRENT APPLICATION NUMBER: US/10/750,185
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 43510
; LENGTH: 1437
; TYPE: DNA
; ORGANISM: Bovine 19866880876041
US-10-750-185-43510

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Query Match          5.0%; Score 34.4; DB 6; Length 1437;
Best Local Similarity 51.3%; Pred. No. 0.61;
Matches 80; Conservative 0; Mismatches 76; Indels 0; Gaps 0;

Qy 104 CTTATGGGGACCCCTGGCCACAGCTGCTCTCTCTTGGCCCTCTTTGGTACAGGGAGG 163
Db 1394 CCTCAAGAGGAGGAGTGCAGGGCTGCCCTTCTCTGCTCTGCTCTGCTCTGCTCTGCT 1335

Qy 164 AGCAGCTGGCCCATCAGCTCCCATCAGAGGTGACAAAGTCCAACTTCCAGCAGCCCTA 223
Db 1334 ACTACCTGGGGCAGCCCTTCTATGCTGTGGCAGCAGGTGACCACTCTCCAGATTCCAGA 1275

Qy 224 TATACCAACCGCACCTTTCATGCTGCTAAGAGGC 259
Db 1274 CACTCCCAAGAGAGCTCTCGTGACGCCCCCGAGAGC 1239

, RESULT 13

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US-10-750-185-31621/c
; Sequence 31621, Application US/10750185
; Publication No. US20050260603A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-2
; CURRENT APPLICATION NUMBER: US/10/750,185
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 31621
; LENGTH: 3044
; TYPE: DNA
; ORGANISM: Bovine 19866881259754
US-10-750-185-31621

Query Match          4.9%; Score 33.8; DB 6; Length 3044;
Best Local Similarity 51.0%; Pred. No. 1.5;
Matches 80; Conservative 0; Mismatches 77; Indels 0; Gaps 0;

Qy 92 TGTGAGCTCTTTCTTATGGGACCCCTGGCCACAGCTGCTCTCTTCTTGGCCCTCTT 151
Db 2287 TGGTGGCACCTTCTTCTCAGTGAGCCCTCGCGTACAGAGGCTACTTACTTCTGGTCAICAG 2228

Qy 152 GGTACAGGAGGAGCAGCTGCGCCCATCAGTCCCATGCGAGGCTTCCAAAGTCCAACTT 211
Db 2227 TGAACAACACAAGAAAGATGTGCACAGCGCCCACTCTCTCTCTCTCTGGGTAGACCTC 2168

Qy 212 CCAGCAGCCCTATATACCAACCGCACCTTCATGCTG 248
Db 2167 TATCCAGCCCATCTTGTCTCAGCATCTCTTCCCTCTG 2131

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RESULT 14
US-10-750-185-43708
; Sequence 43708, Application US/10750185
; Publication No. US20050260603A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-2
; CURRENT APPLICATION NUMBER: US/10/750,185
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 43708
; LENGTH: 2673
; TYPE: DNA
; ORGANISM: Bovine 19866880749669
US-10-750-185-43708

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Query Match          4.7%; Score 32.6; DB 6; Length 2673;
Best Local Similarity 59.3%; Pred. No. 3.4;
Matches 73; Conservative 0; Mismatches 49; Indels 1; Gaps 1;

Qy 76 CCGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGGCCACAGCTGCTCTCC 135

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Db      27  CCGACCTCAGGGGCACTGTGTCTCGTCTGTGGGGCTGCTGTCTCCCCAGCT-CCTCC 85
Qy      136  TTCTCTTGGCCCTCTTGGTACAGGGAGCAGCAGCTGCGCCCATCAGCTCCCACTGCAGGC 195
Db      86  TCCCTGGGGCCAGTCTCTCCAGCCCCGAGCAGGGAGGCCCGGAGGCATGACAGCAGTC 145
Qy      196  TTG 198
Db      146  TGG 148
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RESULT 15
US-10-750-185-50850
; Sequence 50850, Application US/10750185
; Publication No. US20050260603A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-2
; CURRENT APPLICATION NUMBER: US/10750,185
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 50850
; LENGTH: 2126
; TYPE: DNA
; ORGANISM: Bovine 19866881174911
US-10-750-185-50850
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Query Match      4.7%; Score 32.2; DB 6; Length 2126;
Best Local Similarity 49.0%; Pred. No. 4;
Matches 117; Conservative 0; Mismatches 118; Indels 4; Gaps 1;

Qy      441  AGGCTCAGCAACAGGCTAAGCACATGTCAATTGAAGGTGATGACCTGCATATCCAGAGG 500
Db      1471  AGTTCTGAAAAATCTTTAACACAAGTCACATTACAGGTGATGAGGTAAATAGGAAGG 1530
Qy      501  AATGTGCAAAAGCTGAAG-----GACACAGTGAAAAAGCTTGGAGAGAGTGGAGAGATCAA 556
Db      1531  ATGAAGAAGAAATAAAGAATTTACTCTGTGAAGCGAATTTGAGACAGAGGATTTTCAGAT 1590
Qy      557  AGCAATTGGAGAACTGGATTGCTGTTTATGTCTCTGAGAAATGCCTGCATTTGACCAGA 616
Db      1591  ATTATATTATTACTTCTTGGTCTCTTCTCTCATACACACATATGCACATACTTTCATA 1650
Qy      617  GCAAAGCTGAAAAATGAATAACTAACCCCTTTCCCTGCTAGAAATAACAATTAGATGC 675
Db      1651  TTCATGTTTCAGAGAAATAAGAAGCCACCCTGTTGGAGTAGACAGCATGTAAATAAATAC 1709
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OM nucleic - nucleic search, using sw model

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Perfect score: 4797

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Scoring table: IDENTITY NUC

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Searched: 1303057 seqs, 888780828 residues

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Minimum DB seq length: 0

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Post-processing: Minimum Match 0%

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Listing first 45 summaries

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- 8: /cgn2\_6/ptodata/1/ina/RE/COMB.seq.\*
- 9: /cgn2\_6/ptodata/1/ina/backfiles1.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	4797	100.0	4797	3	US-09-419-568F-25
2	4797	100.0	4797	3	US-09-354-243B-25
3	4738.6	98.8	8888	3	US-09-949-016-17185
4	686	14.3	7445	3	US-09-178-973B-8
5	686	14.3	7445	3	US-09-419-568F-8
6	686	14.3	7445	3	US-09-354-243B-8
7	650	13.6	5935	3	US-09-178-973B-17
8	650	13.6	5935	3	US-09-419-568F-29
9	650	13.6	5935	3	US-09-354-243B-29
10	600.6	12.5	601	3	US-09-949-016-190092
11	600.6	12.5	601	3	US-09-949-016-190093
12	258	5.4	690	3	US-09-419-568F-24
13	258	5.4	690	3	US-09-354-243B-24
14	256	5.3	689	3	US-09-949-016-5443
15	244	5.1	1152	3	US-09-870-574-1
16	237	4.9	1191	3	US-09-084-298-1
17	207	4.3	1116	3	US-10-090-365-14
18	207	4.3	1116	3	US-09-728-911-14
19	140	2.9	601	3	US-09-949-016-86957
20	140	2.9	247781	3	US-09-949-016-14193
21	132.2	2.8	87780	3	US-09-949-016-17011
22	132.2	2.8	90428	3	US-09-949-016-12564
23	130.6	2.7	28213	3	US-09-949-016-12738
24	130.6	2.7	28216	3	US-09-949-016-13652

ALIGNMENTS

RESULT 1

US-09-419-568F-25  
; Sequence 25, Application US/09419568F  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors and Uses Thereof  
; FILE OF INVENTION: (TIFs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/419,568F  
; CURRENT FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 25  
; LENGTH: 4797  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-09-419-568F-25

Query Match	100.0%;	Score 4797;	DB 3;	Length 4797;
Best Local Similarity	100.0%;	Pred. No. 0;		
Matches 4797;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
Qy	1	TGCACAGCAGAAATCTTCAGACAGAGTTCTCTTCCCAAGTACACAGTTCCTCGAGTTAG	60	
Db	1	TGCACAGCAGAAATCTTCAGACAGAGTTCTCTTCCCAAGTACACAGTTCCTCGAGTTAG	60	
Qy	61	AATTGCTGCAATGGCGCCCTTCGAGAAATCTGTAGCTCTTCTTATGGGACCTGG	120	
Db	61	AATTGCTGCAATGGCGCCCTTCGAGAAATCTGTAGCTCTTCTTATGGGACCTGG	120	
Qy	121	CCACCAAGTCCCTCTCTTCTGCGCCCTTCGTACAGGAGGAGGAGTCCGCCCCATCA	180	
Db	121	CCACCAAGTCCCTCTCTTCTGCGCCCTTCGTACAGGAGGAGGAGTCCGCCCCATCA	180	
Qy	181	GCTCCCACTCCAGGCTTGCAAGTCCAACTTCCAGCAGCCCTATATCAACCAACCGACCT	240	
Db	181	GCTCCCACTCCAGGCTTGCAAGTCCAACTTCCAGCAGCCCTATATCAACCAACCGACCT	240	
Qy	241	TGATGCTGGTAAAGGAGGTATACATCTCAATCTCTTCTGCTTGGATCTACTTGA	300	
Db	241	TGATGCTGGTAAAGGAGGTATACATCTCAATCTCTTCTGCTTGGATCTACTTGA	300	

Db 241 TCATGCTGGCTAAGGAGGTATACATCTCAATCTCGTCTCTTTCTCGTTGGATCTACTTGA 300  
Qy 301 ATCCAAATAGTCTTAAACITTTCTTCAAGACATCTCTAAGAGCTTTTAGGAACCCACTGT 360  
Db 301 ATCCAAATAGTCTTAAACITTTCTTCAAGACATCTCTAAGAGCTTTTAGGAACCCACTGT 360  
Qy 361 TTATCCCTGAGGTAGATAAATTTCTTGTTTTTTCAAGAGCTCTTTGGGAATCTGGCTTT 420  
Db 361 TTATCCCTGAGGTAGATAAATTTCTTGTTTTTTCAAGAGCTCTTTGGGAATCTGGCTTT 420  
Qy 421 TTTTTTTCTTGAACITTTCTTCCATTTTGGCTTTATAGATACATATGATGAATTTT 480  
Db 421 TTTTTTTCTTGAACITTTCTTCCATTTTGGCTTTATGATACATATGATGAATTTT 480  
Qy 481 CCCAAGAGCGGCATTCAGTAATCCATCTCATGATTTTTTTTTTCTTATGCTCTGTG 540  
Db 481 CCCAAGAGCGGCATTCAGTAATCCATCTCATGATTTTTTTTTTCTTATGCTCTGTG 540  
Qy 541 CATTTGTTCTAACTCATGACACATCTGAATTTCTGCTTTTATGCTTTTATGATGTTGCTCT 600  
Db 541 CATTTGTTCTAACTCATGACACATCTGAATTTCTGCTTTTATGCTTTTATGATGTTGCTCT 600  
Qy 601 GGGGAGCGGATGGGGCACATGCTATGTATAAAATTTTTTTCTATTTGCTCAATGTCC 660  
Db 601 GGGGAGCGGATGGGGCACATGCTATGTATAAAATTTTTTTCTATTTGCTCAATGTCC 660  
Qy 661 AGACCTTAGTCTTTTCTTCTTCCAGGCTAGCTTGCTGATTAACAACACAGACGTTG 720  
Db 661 AGACCTTAGTCTTTTCTTCTTCCAGGCTAGCTTGCTGATTAACAACACAGACGTTG 720  
Qy 721 TCTCATTTGGGAGAAACTGTTTCCAGGAGTCAGTGTAAAGCTTACAGTTGTGACGAACAGG 780  
Db 721 TCTCATTTGGGAGAAACTGTTTCCAGGAGTCAGTGTAAAGCTTACAGTTGTGACGAACAGG 780  
Qy 781 CCGTGTGCGTCCATGGGTACTTGGGGTGGTGGTATGATGGTTTAGTCTTTATCCCTTA 840  
Db 781 CCGTGTGCGTCCATGGGTACTTGGGGTGGTGGTATGATGGTTTAGTCTTTATCCCTTA 840  
Qy 841 TGACCTTCTGTTTCCCTTCCACTGCGAGATGAGTGAGCGCTGCTATCTGATGAAGCAG 900  
Db 841 TGACCTTCTGTTTCCCTTCCACTGCGAGATGAGTGAGCGCTGCTATCTGATGAAGCAG 900  
Qy 901 GTGCTGAACITTCACCTTTGAAGAAGTCTGTTCCCTCAATCTGATAGGTTCCAGCTTAT 960  
Db 901 GTGCTGAACITTCACCTTTGAAGAAGTCTGTTCCCTCAATCTGATAGGTTCCAGCTTAT 960  
Qy 961 ATGCAAGAGGTGGTGGCTTCTTGGCCAGGCTCAGCAACAGGCTAAGCACATGTGTAAGT 1020  
Db 961 ATGCAAGAGGTGGTGGCTTCTTGGCCAGGCTCAGCAACAGGCTAAGCACATGTGTAAGT 1020  
Qy 1021 TCAGCTCTCAGGCTATGCGCCACTTACCCCTCGTTCCCTCTTCCACAGAGACCCCTTAC 1080  
Db 1021 TCAGCTCTCAGGCTATGCGCCACTTACCCCTCGTTCCCTCTTCCACAGAGACCCCTTAC 1080  
Qy 1081 CCCAACTCTCTCTCCCTTCCCTTACCTAGCTAGCAGGAAGTGTCTTGGCAGCAG 1140  
Db 1081 CCCAACTCTCTCTCTCCCTTCCCTTACCTAGCTAGCAGGAAGTGTCTTGGCAGCAG 1140  
Qy 1141 TGTATCAGGAGTCATTTGGGATCATAGATATTTGCTTTTGTGACTCAGTCAATC 1200  
Db 1141 TGTATCAGGAGTCATTTGGGATCATAGATATTTGCTTTTGTGACTGAGTCAATC 1200  
Qy 1201 TTGAGTTATAGTGTGAATGGGTCTGGAATTAAGTGTACAGAAGCCGATTTGGTTTG 1260  
Db 1201 TTGAGTTATAGTGTGAATGGGTCTGGAATTAAGTGTACAGAAGCCGATTTGGTTTG 1260  
Qy 1261 TCTTCGGAAGAAAGCCAACTCAGTGTGGTGAAGTGAAGAGTGTTCGGAAACATCTA 1320  
Db 1261 TCTTCGGAAGAAAGCCAACTCAGTGTGGTGAAGTGAAGAGTGTTCGGAAACATCTA 1320  
Qy 1321 GCTGTGGAATGGATTCATTTAGTCTAAGTTGTTGAGGGGAGGGGATGGCATGGAGAA 1380  
Db 1321 GCTGTGGAATGGATTCATTTAGTCTAAGTTGTTGAGGGGAGGGGATGGCATGGAGAA 1380

Qy 1381 ATTAGAAGAGAAAAGTGGGAAATGGGAAGGCTTAAAGTCGTCGTCGGTTCGCGACACTGTT 1440  
Db 1381 ATTAGAAGAGAAAAGTGGGAAATGGGAAGGCTTAAAGTCGTCGTCGGTTCGCGACACTGTT 1440  
Qy 1441 GCCCTGTTGATGTATGCGGAAGCCACAAAATCGGAGCGGTGTGAACCTGTATGCGCTGAA 1500  
Db 1441 GCCCTGTTGATGTATGCGGAAGCCACAAAATCGGAGCGGTGTGAACCTGTATGCGCTGAA 1500  
Qy 1501 CATTTGAAACTATGAAAAAAGTTTGTGAGTGGGCGCCAGTAAAGAGCCCTTAGGACTT 1560  
Db 1501 CATTTGAAACTATGAAAAAAGTTTGTGAGTGGGCGCCAGTAAAGAGCCCTTAGGACTT 1560  
Qy 1561 ACTGAAGAGGCTTAAATTTTCAATGAGATGTTTATGTATCATATTTCTTGTCTTAAGCATG 1620  
Db 1561 ACTGAAGAGGCTTAAATTTTCAATGAGATGTTTATGTATCATATTTCTTGTCTTAAGCATG 1620  
Qy 1621 CAATTTTCTGAGATACGATTTAGGTTTTATCTTACAGAAATTTGCAATAAATCTACTCCG 1680  
Db 1621 CAATTTTCTGAGATACGATTTAGGTTTTATCTTACAGAAATTTGCAATAAATCTACTCCG 1680  
Qy 1681 CTCTTTCCACAAATGCAAACTCAGTAGGATTTCCCAAGATGAAGAGAGTCTCTTGTGA 1740  
Db 1681 CTCTTTCCACAAATGCAAACTCAGTAGGATTTCCCAAGATGAAGAGAGTCTCTTGTGA 1740  
Qy 1741 AGGAAGTGTGATTTCTGCGCTCCAAGGGAATTTCAAGAGCTCAGGAAATCTTAGGTCA 1800  
Db 1741 AGGAAGTGTGATTTCTGCGCTCCAAGGGAATTTCAAGAGCTCAGGAAATCTTAGGTCA 1800  
Qy 1801 TGTGAAATCTAGGTCAATTTGGGCAAAATTTACTAAGAGCTTTAAATTTCCAGGTCAATGT 1860  
Db 1801 TGTGAAATCTAGGTCAATTTGGGCAAAATTTACTAAGAGCTTTAAATTTCCAGGTCAATGT 1860  
Qy 1861 ACTGTACTCTCAATGGGTGTGAGGTTTCATAAAGTTTTCAGCACAACTTAAGATAGTTATG 1920  
Db 1861 ACTGTACTCTCAATGGGTGTGAGGTTTCATAAAGTTTTCAGCACAACTTAAGATAGTTATG 1920  
Qy 1921 CTTGTTATTTTATAGCATATTTGAAGGTGATGACCTGCATATCCAGAGAAATGTGCAA 1980  
Db 1921 CTTGTTATTTTATAGCATATTTGAAGGTGATGACCTGCATATCCAGAGAAATGTGCAA 1980  
Qy 1981 AAGCTGAAGGACACAGTGAAGAGTGAAGTGAATCTGTCATATGCTCAATGCTAATGCAAT 2040  
Db 1981 AAGCTGAAGGACACAGTGAAGAGTGAAGTGAATCTGTCATATGCTCAATGCTAATGCAAT 2040  
Qy 2041 AGGAGAGCAAAATGTTGTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2100  
Db 2041 AGGAGAGCAAAATGTTGTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2100  
Qy 2101 CTTGATTTCTCTACCCAGGCGATTTACTTTTGTGTCTGTGTATGTAGATATATCTATA 2160  
Db 2101 CTTGATTTCTCTACCCAGGCGATTTACTTTTGTGTCTGTGTATGTAGATATATCTATA 2160  
Qy 2161 TATCTAGATGTCACTTTCCAAATCTTGAATTTGTAGAAATTTCTAGAACTGTTGGGATCT 2220  
Db 2161 TATCTAGATGTCACTTTCCAAATCTTGAATTTGTAGAAATTTCTAGAACTGTTGGGATCT 2220  
Qy 2221 TAGCTTGTCTAGTCAATACCTCAGATTTCTGGGGATGGTCAAGTGGCAGAGATAGGGCTA 2280  
Db 2221 TAGCTTGTCTAGTCAATACCTCAGATTTCTGGGGATGGTCAAGTGGCAGAGATAGGGCTA 2280  
Qy 2281 GAATGCAAGTCTCTGATCCCAAGCCAGCACTTTTCCCGTGGTGTATGATAGATTTAGTTT 2340  
Db 2281 GAATGCAAGTCTCTGATCCCAAGCCAGCACTTTTCCCGTGGTGTATGATAGATTTAGTTT 2340  
Qy 2341 TGGTACCAATTAATTTCTTAGGGAATTTTCCAGATTTCTTATGTACTCATGTAATCTGAAGAAG 2400  
Db 2341 TGGTACCAATTAATTTCTTAGGGAATTTTCCAGATTTCTTATGTACTCATGTAATCTGAAGAAG 2400  
Qy 2401 TACTGTTTAAAAAAGCAAGAAATGCTATGGGCAAAATTTTATTTGAGTCAATTTTGAAGT 2460  
Db 2401 TACTGTTTAAAAAAGCAAGAAATGCTATGGGCAAAATTTTATTTGAGTCAATTTTGAAGT 2460

QY	2461	CATTAAATGCATTGCTTTGAAACTTTGGAAGTAATAAATCAGAACAAATGAGAAAGAGCTGG	2520
Db	2461	CATTAAATGCATTGCTTTGAAACTTTGGAAGTAATAAATCAGAACAAATGAGAAAGAGCTGG	2520
QY	2521	ACTTGCAATATAGGCTAAATTTCTGGAGTAATAACACTTATTTTGAATTAATCATAAATATC	2580
Db	2521	ACTTGCAATATAGGCTAAATTTCTGGAGTAATAACACTTATTTTGAATTAATCATAAATATC	2580
QY	2581	TATCAGATATTTGATTTATAGTTTAAAGCAAGAGCAGACAACCCCGATCTCTTTTATACAG	2640
Db	2581	TATCAGATATTTGATTTATAGTTTAAAGCAAGAGCAGACAACCCCGATCTCTTTTATACAG	2640
QY	2641	GTTCAAATAGAGTAATAAATATATAGTAAGATTTTATATAGTTAAATGGAAGTCTGAAT	2700
Db	2641	GTTCAAATAGAGTAATAAATATATAGTAAGATTTTATATAGTTAAATGGAAGTCTGAAT	2700
QY	2701	GGTAAGCTTTTTTTCTCTCTCTCCATCAAGACCTTCCATCTAGTTTCTTCCCTTCA	2760
Db	2701	GGTAAGCTTTTTTTCTCTCTCTCCATCAAGACCTTCCATCTAGTTTCTTCCCTTCA	2760
QY	2761	CTCCCTCAACAAATCCCTAGGAGCATTTATCCATGGTGGCTGGTGTACATTTCTATAG	2820
Db	2761	CTCCCTCAACAAATCCCTAGGAGCATTTATCCATGGTGGCTGGTGTACATTTCTATAG	2820
QY	2821	TGAATGATACCATCATGTGGCTTATTTGGTGAAGAAACAACAATGGAAGGCTTAGACTA	2880
Db	2821	TGAATGATACCATCATGTGGCTTATTTGGTGAAGAAACAACAATGGAAGGCTTAGACTA	2880
QY	2881	ACAAATGATACCATCATGTGGCTTATTTGGTGAAGAAACAACAATGGAAGGCTTAGACTA	2940
Db	2881	ACAAATGATACCATCATGTGGCTTATTTGGTGAAGAAACAACAATGGAAGGCTTAGACTA	2940
QY	2941	GCAAGCAGGTACAACTAAATCTCAGAAACATGAAGGCTCCAGTTGATGGAATTTTCAGT	3000
Db	2941	GCAAGCAGGTACAACTAAATCTCAGAAACATGAAGGCTCCAGTTGATGGAATTTTCAGT	3000
QY	3001	AACAAGCTTAACTTAAATCCCTTTTCCCTCTTGACTTTTAAAAAAGCGTTCTTC	3060
Db	3001	AACAAGCTTAACTTAAATCCCTTTTCCCTCTTGACTTTTAAAAAAGCGTTCTTC	3060
QY	3061	CTGAGCATCATTTAATGAGTGTGACTCTTCTCTCTTTGATTAATGGAAGCTTTGTAGTT	3120
Db	3061	CTGAGCATCATTTAATGAGTGTGACTCTTCTCTCTTTGATTAATGGAAGCTTTGTAGTT	3120
QY	3121	TTAAATTTGAGAGCCAGTTCTCTTTGTTATAGAACTATTTCTAGACATGAGAGGCTGAA	3180
Db	3121	TTAAATTTGAGAGCCAGTTCTCTTTGTTATAGAACTATTTCTAGACATGAGAGGCTGAA	3180
QY	3181	TGTTAGCATGCCACAGACAAGGCATGCTTTTACACATCTTGCTTAAAAAATTTACTGATTC	3240
Db	3181	TGTTAGCATGCCACAGACAAGGCATGCTTTTACACATCTTGCTTAAAAAATTTACTGATTC	3240
QY	3241	ATCTTGCTTGTTGTTTAAAGAGTGAAGTGTGAGAGAGAGAAATCTCATGTGTGATCTG	3300
Db	3241	ATCTTGCTTGTTGTTTAAAGAGTGAAGTGTGAGAGAGAGAAATCTCATGTGTGATCTG	3300
QY	3301	TGTGATTTTCAAGACCTTTAATCCATTTTGAAGAACTAATTTTCAATTTTGCATGGGTT	3360
Db	3301	TGTGATTTTCAAGACCTTTAATCCATTTTGAAGAACTAATTTTCAATTTTGCATGGGTT	3360
QY	3361	GCCATGTGGAAGAGTGAATATGCTTTTGTGCTAGCTTCCAGAAAGCAGAGGAGGAGA	3420
Db	3361	GCCATGTGGAAGAGTGAATATGCTTTTGTGCTAGCTTCCAGAAAGCAGAGGAGGAGA	3420
QY	3421	GCAATGTGTTTCAAGAAAGATCAACAGAGGAGAAATCTGTGAGAGTGTCTGAAATAGG	3480
Db	3421	GCAATGTGTTTCAAGAAAGATCAACAGAGGAGAAATCTGTGAGAGTGTCTGAAATAGG	3480
QY	3481	GTGGTTTTTGGAGGCAATTAATCCCTCTCTGTTGGGGTAAAAAGCAGAACCGAGGTTGGTA	3540
Db	3481	GTGGTTTTTGGAGGCAATTAATCCCTCTCTGTTGGGGTAAAAAGCAGAACCGAGGTTGGTA	3540
QY	3541	GTAATAATGCATGACAGACAGTAGGGGACGATAAACTTTAAAAATTTCTTTATAGTCTTGGAG	3600

Db	3541	GTAAAAATGCATGACAGACAGTAGGGGACGATAAACTTTAAAAATTTCTTTATAGTCTTGGAG	3600
QY	3601	TCCTTTGAGATAGAAAAAATATCTTTTGGCCCTTATGTCAAAAAGAACTATGGAAGGTGA	3660
Db	3601	TCCTTTGAGATAGAAAAAATATCTTTTGGCCCTTATGTCAAAAAGAACTATGGAAGGTGA	3660
QY	3661	AAGGGCGGAAAGAAAGCAGGAAAAAGCAACATATATATATATAGAGGACAATGGTGACA	3720
Db	3661	AAGGGCGGAAAGAAAGCAGGAAAAAGCAACATATATATATATATAGAGGACAATGGTGACA	3720
QY	3721	AGGTTTTTCTTTGAAATAATGCAATATATAGTAGATTTAGAGGAAATTTCAAGTGGGAATGCTT	3780
Db	3721	AGGTTTTTCTTTGAAATAATGCAATATATAGTAGATTTAGAGGAAATTTCAAGTGGGAATGCTT	3780
QY	3781	TTCACTTTGAAATTTGGGTTTCTCTTCGATTTAAGTTTGGGATCCCTCATCTGCAATTTGACTT	3840
Db	3781	TTCACTTTGAAATTTGGGTTTCTCTTCGATTTAAGTTTGGGATCCCTCATCTGCAATTTGACTT	3840
QY	3841	GGAGAGAGAAAGAAATGAATGTTAGGACCTATATCTGTTTTTCTTANTTAACATAAGCAAGTG	3900
Db	3841	GGAGAGAGAAAGAAATGAATGTTAGGACCTATATCTGTTTTTCTTANTTAACATAAGCAAGTG	3900
QY	3901	GAAGAAGCTTTATTTGGTATTTTTTCCCAAAAGTGAAAACTTTTCTTTTACTGTTGTCA	3960
Db	3901	GAAGAAGCTTTATTTGGTATTTTTTCCCAAAAGTGAAAACTTTTCTTTTACTGTTGTCA	3960
QY	3961	AAAAGGTGGAAAAATAGAAAAAGCCTTAATGTTATTTGGTGAATACATGTTTCAAGTCATTTG	4020
Db	3961	AAAAGGTGGAAAAATAGAAAAAGCCTTAATGTTATTTGGTGAATACATGTTTCAAGTCATTTG	4020
QY	4021	AGTAGAGATGTTTTTAAATCAGGAGTGTCCAATCATTTTGGCTTCCCTGGACCACTTGAAA	4080
Db	4021	AGTAGAGATGTTTTTAAATCAGGAGTGTCCAATCATTTTGGCTTCCCTGGACCACTTGAAA	4080
QY	4081	GAATTTGCTTTGGTACACATAAAATACAGAAACATAGCTGATGAGCTAAAAAGTCCA	4140
Db	4081	GAATTTGCTTTGGTACACATAAAATACAGAAACATAGCTGATGAGCTAAAAAGTCCA	4140
QY	4141	TGCATAAATCTCATCTACTGTTTTTAAAGAAAGTTTATGAATTTCTGTTAGGGTGCAATTCAAAG	4200
Db	4141	TGCATAAATCTCATCTACTGTTTTTAAAGAAAGTTTATGAATTTCTGTTAGGGTGCAATTCAAAG	4200
QY	4201	CTGTCCTGGGCCCATGTGCGGCCCTGTGGGCTGAGGTTGGAGCAAGCTCCTTATAAGTAATC	4260
Db	4201	CTGTCCTGGGCCCATGTGCGGCCCTGTGGGCTGAGGTTGGAGCAAGCTCCTTATAAGTAATC	4260
QY	4261	TGTCATAGATAGTTTTTGGAGCTGCAAAAACAGGCCCAAGGCATTAATGGGTGGCACTCGGGAT	4320
Db	4261	TGTCATAGATAGTTTTTGGAGCTGCAAAAACAGGCCCAAGGCATTAATGGGTGGCACTCGGGAT	4320
QY	4321	CCCCCAGATCCCAGCCTCACTTCAGTCTCCTCTGCTGCTGTTTAAAGAGGGTGGTCAACTC	4380
Db	4321	CCCCCAGATCCCAGCCTCACTTCAGTCTCCTCTGCTGCTGTTTAAAGAGGGTGGTCAACTC	4380
QY	4381	TCGCCCCAGCTTTTAAACAGCTTCATTAGTGTGAGGTGACCTGAAATTTGATGCTGCTG	4440
Db	4381	TCGCCCCAGCTTTTAAACAGCTTCATTAGTGTGAGGTGACCTGAAATTTGATGCTGCTG	4440
QY	4441	GTGGCCTCTCAGTCCAGAGAGCCGTCAATTTTAAAGCTCTTTGGCAAAATCATACAATACTAA	4500
Db	4441	GTGGCCTCTCAGTCCAGAGAGCCGTCAATTTTAAAGCTCTTTGGCAAAATCATACAATACTAA	4500
QY	4501	AGGGATATTAATGATGATTTTACAAATGCTTAAAACTCGGTTTCTGCTCCATCAACC	4560
Db	4501	AGGGATATTAATGATGATTTTACAAATGCTTAAAACTCGGTTTCTGCTCCATCAACC	4560
QY	4561	TAATCTTGCAATTTCTAAATTTGTTTCACTTTAGAAAAACATGGCATAAATGCTCAAAATCTT	4620
Db	4561	TAATCTTGCAATTTCTAAATTTGTTTCACTTTAGAAAAACATGGCATAAATGCTCAAAATCTT	4620
QY	4621	TTGCATTTCTTATTTTCAAGCTTTGGAGAGAGTGGAGAGATCAAGCAATTTGGAGAACTGG	4680

Db 4621 TTGCAATCTTATTTTACAGCTTGAGAGAGTGGAGAGATCAAAGCAATTGGAGAACTGG 4680  
Qy 4681 ATTTGCTGTTTATGCTCTGAGAAATGCCTGCATTTGACAGAGCAAAAGCTGAAAAATGA 4740  
Db 4681 ATTTGCTGTTTATGCTCTGAGAAATGCCTGCATTTGACAGAGCAAAAGCTGAAAAATGA 4740  
Qy 4741 ATAACTAAACCCCTTTCCCTGCTAGAAATAACAATTAGATGCCCAAGCGATTTTT 4797  
Db 4741 ATAACTAAACCCCTTTCCCTGCTAGAAATAACAATTAGATGCCCAAGCGATTTTT 4797

RESULT 2  
US-09-354-243B-25  
; Sequence 25, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Peptides  
; TITLE OF INVENTION: (TIFB)  
; FILE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.1  
; CURRENT APPLICATION NUMBER: US/09/354.243B  
; CURRENT FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 25  
; LENGTH: 4797  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-09-354-243B-25

Query Match 100.0%; Score 4797; DB 3; Length 4797;  
Best Local Similarity 100.0%; Pred. No. 0;  
Matches 4797; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TGCAAGCAGAGATCTTTGAGAAAGGTTCTCTTCCCGAGTCCAGAGTTGCTCGAGTTAG 60  
Db 1 TGCAAGCAGAGATCTTTGAGAAAGGTTCTCTTCCCGAGTCCAGAGTTGCTCGAGTTAG 60  
Qy 61 AATTGCTGCAATGGCGCCCTGAGAAATCTGTGAGCTCTTCTTATGGGACCCCTGG 120  
Db 61 AATTGCTGCAATGGCGCCCTGAGAAATCTGTGAGCTCTTCTTATGGGACCCCTGG 120  
Qy 121 CCACGAGTGGCTCTCTTCTTGGCCCTCTTGGTACAGGAGGAGCAGCTGCGCCCATCA 180  
Db 121 CCACGAGTGGCTCTCTTCTTGGCCCTCTTGGTACAGGAGGAGCAGCTGCGCCCATCA 180  
Qy 181 GCTCCACTGAGGCTTGACAACTTCCAGAGCCCTATATCAACACCGACCT 240  
Db 181 GCTCCACTGAGGCTTGACAACTTCCAGAGCCCTATATCAACACCGACCT 240  
Qy 241 TCATGCTGGCTAAGGAGTATACATCTCAATCTGCTCTTCTGTTGGATCTACTTGA 300  
Db 241 TCATGCTGGCTAAGGAGTATACATCTCAATCTGCTCTTCTGTTGGATCTACTTGA 300  
Qy 301 ATCCAAATAGTTCTTAAACTTTTCTTACAGAGCATCTTAAGAGCTTTTAGGAACCCACTGT 360  
Db 301 ATCCAAATAGTTCTTAAACTTTTCTTACAGAGCATCTTAAGAGCTTTTAGGAACCCACTGT 360  
Qy 361 TTATCCCTGAGGCTAGATAAAATTTTCTGTTTTTTCAGAGCTCTTTGGGAATCTGGCTTT 420  
Db 361 TTATCCCTGAGGCTAGATAAAATTTTCTGTTTTTTCAGAGCTCTTTGGGAATCTGGCTTT 420  
Qy 421 TTTTCTTCTGAACTTCTCTTCCATTTTGGCCCTTTATGATACATATGATGAAATTTT 480  
Db 421 TTTTCTTCTGAACTTCTCTTCCATTTTGGCCCTTTATGATACATATGATGAAATTTT 480  
Qy 481 CCCAAGAGCGCCATTCAGTAATCCATCTGATGATTTTTTTTCTTATGCTCTGCTG 540  
Db 481 CCCAAGAGCGCCATTCAGTAATCCATCTGATGATTTTTTTTCTTATGCTCTGCTG 540

Db 481 CCCAAGAGCGCCATTCAGTAATCCATCTGATGATTTTTTTTCTTATGCTCTGCTG 540  
Qy 541 CATTTGCTTAAATCTCATGACACATCTGAATTTCTGTTTTTAGTCTTTTATGATGTTGCTCT 600  
Db 541 CATTTGCTTAAATCTCATGACACATCTGAATTTCTGTTTTTAGTCTTTTATGATGTTGCTCT 600  
Qy 601 GGGAGACGGGATGGGGACATCTCTATGTATATAAATTTTTTTTTTCTATTTGCTCAATGCTCC 660  
Db 601 GGGAGACGGGATGGGGACATCTCTATGTATATAAATTTTTTTTTTCTATTTGCTCAATGCTCC 660  
Qy 661 AGACCCCTTAGTCTTTTCTTCCAGGCTAGCTTGGCTGTGATAAACAACAGACGTTG 720  
Db 661 AGACCCCTTAGTCTTTTCTTCCAGGCTAGCTTGGCTGTGATAAACAACAGACGTTG 720  
Qy 721 TCTCATTTGGGAGAACTGTTCCACGGAGTCAGTGAAGCTACAGTTGTGACGAACAGGG 780  
Db 721 TCTCATTTGGGAGAACTGTTTCCACGGAGTCAGTGAAGCTACAGTTGTGACGAACAGGG 780  
Qy 781 CCGTGTCCCTTCCATGGGTACTTTGGGGTGGTGTGATGATGTTTAGGTCTTATCCCTTA 840  
Db 781 CCGTGTCCCTTCCATGGGTACTTTGGGGTGGTGTGATGATGTTTAGGTCTTATCCCTTA 840  
Qy 841 TGACCCCTTTCTGTTTCCCTTCCACCTGACAGATGAGTGAGCGCTCTATCTGATGAACGAG 900  
Db 841 TGACCCCTTTCTGTTTCCCTTCCACCTGACAGATGAGTGAGCGCTCTATCTGATGAACGAG 900  
Qy 901 GTGCTGAACCTTCAACCTTTGAAGAGTCTGTTCCCTCAATCTGATAGTTTCCAGCCCTTAT 960  
Db 901 GTGCTGAACCTTCAACCTTTGAAGAGTCTGTTCCCTCAATCTGATAGTTTCCAGCCCTTAT 960  
Qy 961 ATGCAGAGGTGGTGGCCCTTCTTGGCCAGGCTCAGCAACAGGCTTAAGCACAATGTGTAAGT 1020  
Db 961 ATGCAGAGGTGGTGGCCCTTCTTGGCCAGGCTCAGCAACAGGCTTAAGCACAATGTGTAAGT 1020  
Qy 1021 TCAGCTCTAGCCCTATGCCCACTACCCCTCTTCTTCCCTTCCACAGAGACCCCTTAC 1080  
Db 1021 TCAGCTCTAGCCCTATGCCCACTACCCCTCTTCTTCCCTTCCACAGAGACCCCTTAC 1080  
Qy 1081 CCCAACTCTCTCTCTTCCCTTACCCCTAAGCTAGCAGGAGAGAGTGTCTTGGCAGCAG 1140  
Db 1081 CCCAACTCTCTCTCTTCCCTTACCCCTAAGCTAGCAGGAGAGAGTGTCTTGGCAGCAG 1140  
Qy 1141 TGTATCAGAGTCAATTTGGGATCATAGAGTATTTGCTTTTGTCTTGTGACTGAGTCAATC 1200  
Db 1141 TGTATCAGAGTCAATTTGGGATCATAGAGTATTTGCTTTTGTGACTGAGTCAATC 1200  
Qy 1201 TTGAGTTTATAGTGGTGAATGGGCTCTGGAATCTTAAGTGTACAGAAGCGCATTTGGTTG 1260  
Db 1201 TTGAGTTTATAGTGGTGAATGGGCTCTGGAATCTTAAGTGTACAGAAGCGCATTTGGTTG 1260  
Qy 1261 TCTTCGAAAAAAGGCAACTCAGGTTGCTAAGATGAGAAAGGTTTGGGAAAAACATCTA 1320  
Db 1261 TCTTCGAAAAAAGGCAACTCAGGTTGCTAAGATGAGAAAGGTTTGGGAAAAACATCTA 1320  
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Db 1321 GCTGTGAAATGGATCCATTTGAGTCTAAGTTGTTGAGGGAGGGGATGGCATGGAGAGAA 1380  
Qy 1381 ATTAGAAGAGAAAGTGGGAAATGGGAAGGCTTAAAGTCGGTGGTGGGTCGGCAGACTGT 1440  
Db 1381 ATTAGAAGAGAAAGTGGGAAATGGGAAGGCTTAAAGTCGGTGGTGGGTCGGCAGACTGT 1440  
Qy 1441 GCCCTGTGATGTCATGGGAAGCCACAATTCGAGGCGGTGTGAATTTGATGCCGCTGAA 1500  
Db 1441 GCCCTGTGATGTCATGGGAAGCCACAATTCGAGGCGGTGTGAATTTGATGCCGCTGAA 1500  
Qy 1501 CATTTGAACTATCAAAAAGTTTGTAGTGAGTGGGCCCAAGTAAAGGCCCTTAGGACTT 1560  
Db 1501 CATTTGAACTATCAAAAAGTTTGTAGTGAGTGGGCCCAAGTAAAGGCCCTTAGGACTT 1560  
Qy 1561 ACTGAAGAGGGCTTAAATTTTTCACATGAGATGTTTTTATGATCTTCTTGTCTTAAGCATG 1620  
Db 1561 ACTGAAGAGGGCTTAAATTTTTCACATGAGATGTTTTTATGATCTTCTTGTCTTAAGCATG 1620

[illegible]

Qy	2701	GGTAAGCTTTTCTTCTCTCCCATCAAGACCTTCCATCTCTAGTTTCTTCCCTTCA	2760
Db	2701	GGTAAGCTTTTCTTCTCTCCCATCAAGACCTTCCATCTCTAGTTTCTTCCCTTCA	2760
Qy	2761	CTCCCTCAACAAATCCCTAGGGAGCATTTATCCATGCTGGGCTGGTGATACATTTCTATAG	2820
Db	2761	CTCCCTCAACAAATCCCTAGGGAGCATTTATCCATGCTGGGCTGGTGATACATTTCTATAG	2820
Qy	2821	TGAATGATACCATCATGTGTGGCCTATTTTGGTGAAAAAGAAACAACATCGAAGGCTTAGACTA	2880
Db	2821	TGAATGATACCATCATGTGTGGCCTATTTTGGTGAAAAAGAAACAACATCGAAGGCTTAGACTA	2880
Qy	2881	ACAATAGTACTCACCCCAAAAACCGGAGGAAATGATTTAGGAGCAGTGAAAAGTGCAGCTCTT	2940
Db	2881	ACAATAGTACTCACCCCAAAAACCGGAGGAAATGATTTAGGAGCAGTGAAAAGTGCAGCTCTT	2940
Qy	2941	GCAAGCAGGTACAACTAAATPACTCAGAAACATGAAGGCTCCAGTTGATGGAATTTTCAGT	3000
Db	2941	GCAAGCAGGTACAACTAAATPACTCAGAAACATGAAGGCTCCAGTTGATGGAATTTTCAGT	3000
Qy	3001	AACAAGCTTAACCTTAATTCCCTCTTTTCCCTCTTTGACTTTTTAAAAAGGGTTTCTTC	3060
Db	3001	AACAAGCTTAACCTTAATTCCCTCTTTTCCCTCTTTGACTTTTTAAAAAGGGTTTCTTC	3060
Qy	3061	CTGAGCATCAATTAATGAGGTGACTGTGTTCTTCTCTTGTGANTATGGAAGGCTTTGAGTT	3120
Db	3061	CTGAGCATCAATTAATGAGGTGACTGTGTTCTTCTTGTGANTATGGAAGGCTTTGAGTT	3120
Qy	3121	TTAAATTGTGAAGCCAGCTCTCTGTTATAGAACTATTATCTAGACATGGAAGGCTGAA	3180
Db	3121	TTAAATTGTGAAGCCAGCTCTCTGTTATAGAACTATTATCTAGACATGGAAGGCTGAA	3180
Qy	3181	TGTTAGCATGCCACAGACAGGCATGCTTTTACACATCTTTGCTTTAAAAAATTACTGATTTT	3240
Db	3181	TGTTAGCATGCCACAGACAGGCATGCTTTTACACATCTTTGCTTTAAAAAATTACTGATTTT	3240
Qy	3241	ATCTTGCTGTTGCTTTTAGAAAAGTGAAGTGAGAGAGGAGAACTCTCATGTGATCTG	3300
Db	3241	ATCTTGCTGTTGCTTTTAGAAAAGTGAAGTGAGAGAGGAGAACTCTCATGTGATCTG	3300
Qy	3301	TGTGATTTTCAAGACCTTTAATCCATTTTGAAGATCAATTTCATATTTGCAATGGGTT	3360
Db	3301	TGTGATTTTCAAGACCTTTAATCCATTTTGAAGATCAATTTCATATTTGCAATGGGTT	3360
Qy	3361	GCCATGTGAAGAGTGATATGCTTTTGTGCTGTTAGCTTCAGAAAGCACAGAGGGAGA	3420
Db	3361	GCCATGTGAAGAGTGATATGCTTTTGTGCTGTTAGCTTCAGAAAGCACAGAGGGAGA	3420
Qy	3421	GCAATGTTGTTCAGAGAAAGATCAACAGGAGGAGAAACTGTCTGTAATAGG	3480
Db	3421	GCAATGTTGTTCAGAGAAAGATCAACAGGAGGAGAAACTGTCTGTAATAGG	3480
Qy	3481	GTGGTTTGGAGGCAATTAATCCCTCTCGTTGGGGTAAAGCAGAAACGAGGTTGGTA	3540
Db	3481	GTGGTTTGGAGGCAATTAATCCCTCTCGTTGGGGTAAAGCAGAAACGAGGTTGGTA	3540
Qy	3541	GTAATATGCATGACAGACAGTAGGGACGATAACTTTTAAAAATTTCTTTATAGTCTTTGGAG	3600
Db	3541	GTAATATGCATGACAGACAGTAGGGACGATAACTTTTAAAAATTTCTTTATAGTCTTTGGAG	3600
Qy	3601	TCTTTGAGATAGAAAAAGAAATATCTTTTGGCCTTATGTCAAAAGAAAGTATGGAAGGTTGA	3660
Db	3601	TCTTTGAGATAGAAAAAGAAATATCTTTTGGCCTTATGTCAAAAGAAAGTATGGAAGGTTGA	3660
Qy	3661	AAGGGCGGAAAGACAGGAAAGGAAGAACCATGTTATATAGAGGACATTTGAGGAA	3720
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Qy	3721	AGGTTTTCTTGTGAAATAATGCAAAATGATAGATTAGAGGAATTTTCAGTAGGAAATGCTT	3780
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Qy	3781	TTCACTTGAATTTTGGGTTTCTCTTCGATTAAGTTTGGGATCCTCATCTGCAATTTGACTT	3840



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Db 3781 TTCACTTGAATTTGGGTTTCTCTTCGATTAAAGTTTGGGATCCTCATCTGCAATTGACTT 3840
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Qy 4081 GAATTTGCTTGGTACACATATAAATAACAAGCAATAGCTGATGAGCTAAAGTCCA 4140
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Db 4201 CTGTCTTGGGCCATGTGGGGCTGTGGGCTGCAGGTTGGACAAGCTCCTTATAAGTAATC 4260
Qy 4261 TGTATAGATAGTTTGGAGCTGCAAAACAGCCCAAGGCAATGAGGTTGGGCTCGGGAT 4320
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Db 4561 TAATCTGCAATTTCTAATTTGTTTACATTTAGAAAACATGAGCAATAATGCTCAAACTTT 4620
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Db 4681 ATTTGCTGTTTATGCTCTGGAATGCTTCAATTTGACAGCAAAAGCTGAAAAATGA 4740
Qy 4741 ATAACTAACCCCTTTCCCTGCTAGAAAATAACAATAGATGCCCAAGCGATTTT 4797
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RESULT 3  
US-09-949-016-17185  
; Sequence 17185, Application US/09949016  
; Patent No. 6812339  
; GENERAL INFORMATION:

; APPLICANT: VENTER, J. Craig et al.  
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED  
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF  
; FILE REFERENCE: CL001307  
; CURRENT APPLICATION NUMBER: US/09/949,016  
; CURRENT FILING DATE: 2000-04-14  
; PRIOR APPLICATION NUMBER: 60/241,755  
; PRIOR FILING DATE: 2000-10-20  
; PRIOR APPLICATION NUMBER: 60/237,768  
; PRIOR FILING DATE: 2000-10-03  
; PRIOR APPLICATION NUMBER: 60/231,498  
; PRIOR FILING DATE: 2000-09-08  
; NUMBER OF SEQ ID NOS: 207012  
; SOFTWARE: Fast-Seq for Windows Version 4.0  
; SEQ ID NO 17185  
; LENGTH: 8888  
; TYPE: DNA  
; ORGANISM: Human  
US-09-949-016-17185

Query Match 98.8%; Score 4738.6; DB 3; Length 8888;  
Best Local Similarity 99.9%; Pred. No. 0;  
Matches 4774; Conservative 0; Mismatches 4; Indels 3; Gaps 3;

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Qy 80 CTTGCAGAAATCTGTAGCTCTTCTTATGGGACCTGGCCACCAAGCTGCTCTTCT 139
Db 2167 CTTGCAGAAATCTGTAGCTCTTCTTATGGGACCTGGCCACCAAGCTGCTCTTCT 2226
Qy 140 CTTGGCCCTCTTGTTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGCAAGCTTGA 199
Db 2227 CTTGGCCCTCTTGTTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGCAAGCTTGA 2286
Qy 200 CAAGTCCAACCTTCAGAGCCCTTATATCACAAACCGCACCTTCAATGCTGGCTAAGGAGGT 259
Db 2287 CAAGTCCAACCTTCAGAGCCCTTATATCACAAACCGCACCTTCAATGCTGGCTAAGGAGGT 2346
Qy 260 ATACATCTCAATCTGCTCTTCTGTTGATCTACTTGGAAATCCAATAGTTCTTAAAC 319
Db 2347 ATACATCTCAATCTGCTCTTCTGTTGATCTACTTGGAAATCCAATAGTTCTTAAAC 2406
Qy 320 TTTTCTTCAGAGCATCTTAAGAGCTTTAGGAACCCACTGTTTATCCTGAGGGTAGATA 379
Db 2407 TTTTCTTCAGAGCATCTTAAGAGCTTTAGGAACCCACTGTTTATCCTGAGGGTAGATA 2466
Qy 380 AATTTTCTGTTTTTTCAGAGACTCTTTGGGAATCTGGC-TTTTTTTTTTCTTGAACCTC 438
Db 2467 AATTTTCTGTTTTTTCAGAGACTCTTTGGGAATCTGGC-TTTTTTTTTTCTTGAACCTC 2526
Qy 439 TTCTTTCATTTTGGCCCTTTATGATACATATGATGAATTTTCCCAAGAGCGGCCATTC 498
Db 2527 TTCTTTCATTTTGGCCCTTTATGATACATATGATGAATTTTCCCAAGAGCGGCCATTC 2586
Qy 499 AGTAATCCATCTGATGA-TTTTTTTTTTCTTATGCTCTCTGCAATGTTCTTAAACTCAT 557
Db 2587 AGTAATCCATCTGATGAATTTTTTTTTTCTTATGCTCTCTGCAATGTTCTTAAACTCAT 2646
Qy 558 GCACACATCTGAATTTCTGCTTTTATGATGTTTGTCTCTGGGGAGACGGGATGGG 617
Db 2647 GCACACATCTGAATTTCTGCTTTTATGATGTTTGTCTCTGGGGAGACGGGATGGG 2706
Qy 618 CACATGCTATGATAAATTTTTTTTCTTATTTGCTCAATGCTCCAGACCTTAGTCTTTTC 677
Db 2707 CACATGCTATGATAAATTTTTTTTCTTATTTGCTCAATGCTCCAGACCTTAGTCTTTTC 2766
Qy 678 TTCTCTCCAGGCTAGCTTGGCTGATACACACAGAGCTTCTGCTCAATTTGGGGAGAAAC 737
Db 2767 TTCTCTCCAGGCTAGCTTGGCTGATACACACAGAGCTTCTGCTCAATTTGGGGAGAAAC 2826
Qy 738 TGTTCACGGAGTCAGTGTAAAGCTACAGTTGTGACGAAACAGGGCGGTGTGCGCTCCATGG 797
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Db		2827	TGTTCCAGGAGTCAGTGTAAAGCTACAGTTGTGACGAACGGGGCCGTGTCCGTCCTATGG	2886
Qy		798	GTACTTGGGGTGGTGTAGTATGATGGTTTAGTCTCTTATCCCTTATGACCCCTTCTGTTTCC	857
Db		2887	GTACTTGGGGTGGTGTAGTATGATGGTTTAGTCTCTTATCCCTTATGACCCCTTCTGTTTCC	2946
Qy		858	CTTCCACTGTCAGATGAGTGGCCCTGTATCTCATGAAGCAGGAGTGTGAACTTCAACCT	917
Db		2947	CTTCCACTGTCAGATGAGTGGCCCTGTATCTCATGAAGCAGGAGTGTGAACTTCAACCT	3006
Qy		918	TGAAGAGTGTCTGTTCCCTCAATCTGATAGTTCAGGCTTATATGACGAGGTGGTGC	977
Db		3007	TGAAGAGTGTCTGTTCCCTCAATCTGATAGTTCAGGCTTATATGACGAGGTGGTGC	3066
Qy		978	CTTCTCGCCAGGCTCAGCAACAGGCTAAGCACATGTGTAAAGTTTCAAGCTTCTCAGCCTATG	1037
Db		3067	CTTCTCGCCAGGCTCAGCAACAGGCTAAGCACATGTGTAAAGTTTCAAGCTTCTCAGCCTATG	3126
Qy		1038	CCCACTTACCCCTCTCTTCCCTCTTCCACAGAGACCCCTTACCCCAACTCTCTCTCTT	1097
Db		3127	CCCACTTACCCCTCTCTTCCCTCTTCCACAGAGACCCCTTACCCCAACTCTCTCTCTT	3186
Qy		1098	CCCCCTACCCCTTAGCTAGCAGGAAGAGTGTCTTGGCAGCAGTGTATCAGGAGTCATT	1157
Db		3187	CCCCCTACCCCTTAGCTAGCAGGAAGAGTGTCTTGGCAGCAGTGTATCAGGAGTCATT	3246
Qy		1158	TGGGATCATAGATATTTGCTTTGCTTGTACTCAGTTCACATCTTGAGTTTATAGTGGTG	1217
Db		3247	TGGGATCATAGATATTTGCTTTGCTTGTACTCAGTTCACATCTTGAGTTTATAGTGGTG	3306
Qy		1218	AATGGGCTCTGGAACTTAAAGTGTACAGAAGCCGCAATTTGGTTGTCTTCGGAAGAAAGGCA	1277
Db		3307	AATGGGCTCTGGAACTTAAAGTGTACAGAAGCCGCAATTTGGTTGTCTTCGGAAGAAAGGCA	3366
Qy		1278	ACTCAGGTTGCGTAAAGTGTAGAAAGGTGTTGGG-AAAAATCTAGCTGTGGAATGGAATC	1336
Db		3367	ACTCAGGTTGCGTAAAGTGTAGAAAGGTGTTGGGAAAAAACAATCTAGATGTGGNAATGGATC	3426
Qy		1337	CATTGAGTCTAAGTTGTTAGGGGAGGGGATGGCATGGAGAGAAATTAGAAGAGAAAGTG	1396
Db		3427	CATTGAGTCTAAGTTGTTAGGGGAGGGGATGGCATGGAGAGAAATTAGAAGAGAAAGTG	3486
Qy		1397	GGAAATGGGAAGGCTTAAAGTTCGGTGGGTGGGTCGGCAGACTGTTGCCCTGTGATGTGAT	1456
Db		3487	GGAAATGGGAAGGCTTAAAGTTCGGTGGGTGGGTCGGCAGACTGTTGCCCTGTGATGTGAT	3546
Qy		1457	GGGAAGCCACAAAATCGAGGCGTGTGAACCTTGATGCGCTGAAACATTTGAAACTATGAA	1516
Db		3547	GGGAAGCCACAAAATCGAGGCGTGTGAACCTTGATGCGCTGAAACATTTGAAACTATGAA	3606
Qy		1517	AAAAAGTTTCAGTGGAGTGGGCCAGTAAAGGCCCTAGGACTTACTGAAAGAGGCTTAA	1576
Db		3607	AAAAAGTTTCAGTGGAGTGGGCCAGTAAAGGCCCTAGGACTTACTGAAAGAGGCTTAA	3666
Qy		1577	TTTTTCAATGAGATGTTTTATGTACATTTCTTGTCTAAGCATGCAATTTTCTGGAGATA	1636
Db		3667	TTTTTCAATGAGATGTTTTATGTACATTTCTTGTCTAAGCATGCAATTTTCTGGAGATA	3726
Qy		1637	CGATTGAGGTTTTATTCCTTACAGAAATTTGATTAACATCTCCGCTCTTTCCACAAATGC	1696
Db		3727	CGATTGAGGTTTTATTCCTTACAGAAATTTGATTAACATCTCCGCTCTTTCCACAAATGC	3786
Qy		1697	AAACCTCAGTAGGATTTCCCAAGATGACAGAGGTCTCTTGTAAAGGAGTGTGACTGGAT	1756
Db		3787	AAACCTCAGTAGGATTTCCCAAGATGACAGAGGTCTCTTGTAAAGGAGTGTGACTGGAT	3846
Qy		1757	TCTGGCGTCCAAAGGGAATTCAGAGCTCAGGAATCTAGGTCACTGTTGAAATCTAGGTC	1816
Db		3847	TCTGGCGTCCAAAGGGAATTCAGAGCTCAGGAATCTAGGTCACTGTTGAAATCTAGGTC	3906
Qy		1817	ATTGTGGGCAAAATTAATAAGCTTTTAAATCCAGGTGAAATGTACTGTACCTCCATGGG	1876

Db		3907	ATTGTGGGCAAAATTAATAAGAGCTTTTAATTCAGGTGAATGTACTGTACCTCCATGGG	3966
Qy		1877	TGTGGAGTTTCATAAAGTTTTCAGCAACAATAAGATAGTATGCTTGTATTGTTTTAT	1936
Db		3967	TGTGGAGTTTCATAAAGTTTTCAGCAACAATAAGATAGTATGCTTGTATTGTTTTAT	4026
Qy		1937	AGCATATTGAAGGTGATGACCTGTGATATCCAGAGGAATGTCAAAAGCTGAAGACACAG	1996
Db		4027	AGCATATTGAAGGTGATGACCTGTGATATCCAGAGGAATGTCAAAAGCTGAAGACACAG	4086
Qy		1997	TGAAAAGGTGAGGACTGATAAATCTCAATCTAAGTCAATCAATAGGAGACAAAATGTT	2056
Db		4087	TGAAAAGGTGAGGACTGATAAATCTCAATCTAAGTCAATCAATAGGAGACAAAATGTT	4146
Qy		2057	GTTTTTCTTTCTTTCTTTCTTTCCCATCACTTTTGTGATTTTTTCACTTGTATTCTCTACCA	2116
Db		4147	GTTTTTCTTTCTTTCTTTCTTTCCCATCACTTTTGTGATTTTTTCACTTGTATTCTCTACCA	4206
Qy		2117	CCAGGGCGATTACTTTTGGTGTCTGTGTATGTAGATATATCTATATATCTAGATGTCAAGTT	2176
Db		4207	CCAGGGCGATTACTTTTGGTGTCTGTGTATGTAGATATATCTATATATCTAGATGTCAAGTT	4266
Qy		2177	TCCAAATCTTGCAAAATTTAGAAATCTTAGAACTGGTTGGGATCTTAGCTTGTCTAGTCAC	2236
Db		4267	TCCAAATCTTGCAAAATTTAGAAATCTTAGAACTGGTTGGGATCTTAGCTTGTCTAGTCAC	4326
Qy		2237	ATAACCTCAGATTCTGGGGATGGTCAGTGGCAGAGATAGGGCTAGAAATGCAAGTCTCCTG	2296
Db		4327	ATAACCTCAGATTCTGGGGATGGTCAGTGGCAGAGATAGGGCTAGAAATGCAAGTCTCCTG	4386
Qy		2297	AATCCCAAGCCAGACATTTTCCCGTGGTGATACAGATTAAGTTTGGTGTACATTAATCT	2356
Db		4387	AATCCCAAGCCAGACATTTTCCCGTGGTGATACAGATTAAGTTTGGTGTACATTAATCT	4446
Qy		2357	TAGGGAATTTTCAGATTCCTATTTGACTCATGTAATCTGGAAGAGTACTCTGTTTAAACA	2416
Db		4447	TAGGGAATTTTCAGATTCCTATTTGACTCATGTAATCTGGAAGAGTACTCTGTTTAAACA	4506
Qy		2417	GAATAATGCCATATGGGCAAAATTTTATTTGAAAGTCAATTTTGAAGTCAATTAATGCTT	2476
Db		4507	GAATAATGCCATATGGGCAAAATTTTATTTGAAAGTCAATTTTGAAGTCAATTAATGCTT	4566
Qy		2477	TGAAATCTTGGAAGATAAATCTAGAAACAATAGAAAAGAGCTGGACTTGCATATAGGGCT	2536
Db		4567	TGAAATCTTGGAAGATAAATCTAGAAACAATAGAAAAGAGCTGGACTTGCATATAGGGCT	4626
Qy		2537	AATTTCTGGAGTAAATAACACTTTTGTGAAATTTATCATATATCTATCAGATTTGATTA	2596
Db		4627	AATTTCTGGAGTAAATAACACTTTTGTGAAATTTATCATATATCTATCAGATTTGATTA	4686
Qy		2597	TAGTTTAAAGCAAGAGCAGACAAACCCGATCTCTTTTATACAGGTTCAAAATAGAGTAA	2656
Db		4687	TAGTTTAAAGCAAGAGCAGACAAACCCGATCTCTTTTATACAGGTTCAAAATAGAGTAA	4746
Qy		2657	AATATTAGTAAGAGATTTATATTAGTTAAATGGAAGTCTGAAATGGTGAAGCTTTTTTTC	2716
Db		4747	AATATTAGTAAGAGATTTATATTAGTTAAATGGAAGTCTGAAATGGTGAAGCTTTTTTTC	4806
Qy		2717	TTCTCTCTCCCATCAAGACCTTCCATTCTAGTTTCTTCTTCTTCTTCTTCTTCTTCTTCT	2776
Db		4807	TTCTCTCTCCCATCAAGACCTTCCATTCTAGTTTCTTCTTCTTCTTCTTCTTCTTCTTCT	4866
Qy		2777	CTAGGAGCAATTTTATCCATGGTGGGCTGGGTACATTTCTATAGTGAATGATCCATCAT	2836
Db		4867	CTAGGAGCAATTTTATCCATGGTGGGCTGGGTACATTTCTATAGTGAATGATCCATCAT	4926
Qy		2837	GTGGCCTATTGTGAAAAAACAACAATGGAAGGCTTAGACTTAAACAATAGTCACTCACC	2896
Db		4927	GTGGCCTATTGTGAAAAAACAACAATGGAAGGCTTAGACTTAAACAATAGTCACTCACC	4986
Qy		2897	CCAAAACCGGAGGAATTTAGGAGCAATGGAAGTGAAGCTCTTCTGCAAGCAGGTACAACT	2956
Db		4987	CCAAAACCGGAGGAATTTAGGAGCAATGGAAGTGAAGCTCTTCTGCAAGCAGGTACAACT	5046

QY 2957 AAATACTCAGAAACATGAAGCTCCAGTTCATGGAATTTTTCAGTAACAAGCTTAACCTTA 3016  
DB  
QY 5047 AAATACTCAGAAACATGAAGCTCCAGTTCATGGAATTTTTCAGTAACAAGCTTAACCTTA 5106  
DB  
QY 3017 ATTCCTCCCTTTTCCCTCTTGACCTTTTAAAGAGCGTTTCTTCTCCTGAGCATCATTTAAT 3076  
DB  
QY 5107 ATTCCTCCCTTTTCCCTCTTGACCTTTTAAAGAGCGTTTCTTCTCCTGAGCATCATTTAAT 5166  
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DB  
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DB  
QY 5227 AGTTCTCTTGTATAGAACTATTTATCTAGACATGAGGCGTGAATGTTAGCATGCCACAG 5286  
QY 3197 ACAAGGCATGCTTTTACACATCTTCTTAAAGAAATTTACTGATTTTCATCTTGTCTTGTCT 3256  
DB  
QY 5287 ACAAGGCATGCTTTTACACATCTTCTTAAAGAAATTTACTGATTTTCATCTTGTCTTGTCT 5346  
QY 3257 TTAGAAAGTGAAGTGTGAGAGAGAGAACTCTCATGTTGATCTGTGTGATTTTCAAGACC 3316  
DB  
QY 5347 TTAGAAAGTGAAGTGTGAGAGAGAGAACTCTCATGTTGATCTGTGTGATTTTCAAGACC 5406  
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QY 5407 TTTAATCCATTTTGAAGAAATCAATTTTCATATTTGCAATGGGTTGCCATGTGGAGAGTG 5466  
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DB  
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DB  
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DB  
QY 5707 GAATATCTTTTGGCCTTATGTCAAAAGAGATGTGAAAAGTGAAGGCGGAGAAAGC 5766  
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DB  
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DB  
QY 5827 AATGCAAAATATGATAGATTAGAGGAATTTTCAGTAGGGAATGCTTTTCACTTGAATTTGGG 5886  
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DB  
QY 5887 TTTCTCTCTTCGATTAAGTTTGGATTCCTCATCTGCAATTTGACTTGGAGAGAGAAAGATG 5946  
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DB  
QY 5947 AATGTTAGACCTATATCTGTTTCTATTAATTAAGCAAGTGAAGAGACTTATTTGG 6006  
QY 3917 TATTTTCCCAAAAGTGAAACCTTTTCTTTTACTGTTTGTCAAAAGGTGGAATAGA 3976  
DB  
QY 6007 TATTTTCCCAAAAGTGAAACCTTTTCTTTTACTGTTTGTCAAAAGGTGGAATAGA 6066  
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QY 4037 ATCAGGAGTGTCCAATCAATTTGGCTTCCCTGGACCACTTTGAAAGAAATTTCTTTGGTACA 4096  
DB  
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DB  
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QY 4157 TGTTTTAAAGAAAGCTTTATGAATTTCTGTTAGGGTGCATTCAAAAGCTGCTCCTGGGCCATGT 4216  
DB  
QY 6247 TGTTTTAAAGAAAGCTTTATGAATTTCTGTTAGGGTGCATTCAAAAGCTGCTCCTGGGCCATGT 6306  
QY 4217 GCGCCCTGTGGGCTGCAGGTTGGACAAAGCTCCTTATAAGTAAATCTGTGCATAGATGTTTT 6366  
DB  
QY 4277 GAGCTTGCAAAAACAGGCCAAAGGCATTAATGGGTGGCACTCGGGATCCCCAGATCCCCAGCC 4336  
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DB  
QY 6427 TCACCTTCAGTCTCCTTCTGCTGTTAAGAGGGGTGGTCAACTCTCTGCCCCAGCTTTTAA 6486  
QY 4397 ACAGCTTCATTAAGTGTGAGGTGCACCTGAAATTTGATGCCCTGCTGGTGGCCTCTCAGTCCA 4456  
DB  
QY 6487 ACAGCTTCATTAAGTGTGAGGTGCACCTGAAATTTGATGCCCTGCTGGTGGCCTCTCAGTCCA 6546  
QY 4457 GAGAGCGCTCATTTTAAGTCTTTTGGCAATTCATACAAATCTAAAGGGATATTTACTATGA 4516  
DB  
QY 6547 GAGAGCGCTCATTTTAAGTCTTTTGGCAATTCATACAAATCTAAAGGGATATTTACTATGA 6606  
QY 4517 ATGTTTTACAAATGCTTTAAACCTCGGTTCTGCTCTCCATCAACCTTAATCTTTGCAATTTCT 4576  
DB  
QY 6607 ATGTTTTACAAATGCTTTAAACCTCGGTTCTGCTCTCCATCAACCTTAATCTTTGCAATTTCT 6666  
QY 4577 AATTTGTTCACTTTTAGAAAAACATGGCATAAATGCTCAAAATCTTTTGCATTTCTTTATTTTC 4636  
DB  
QY 6667 AATTTGTTCACTTTTAGAAAAACATGGCATAAATGCTCAAAATCTTTTGCATTTCTTTATTTTC 6726  
QY 4637 ACAGCTTGGAGAGAGTGGAGAGATCAAAAGCAATTTGGAGAACTGGATTTGCTGTTTATGTC 4696  
DB  
QY 6727 ACAGCTTGGAGAGAGTGGAGAGATCAAAAGCAATTTGGAGAACTGGATTTGCTGTTTATGTC 6786  
QY 4697 TCTGAGAAATCCCTGCAATTTGACCAGAGCAAGCTGAAAAATGAATAACTAACCCCTTTT 4756  
DB  
QY 6787 TCTGAGAAATCCCTGCAATTTGACCAGAGCAAGCTGAAAAATGAATAACTAACCCCTTTT 6846  
QY 4757 CCCTGCTAGAAATAACAAATTAGATGCCCCAAAGCGATTTTTT 4797  
DB  
QY 6847 CCCTGCTAGAAATAACAAATTAGATGCCCCAAAGCGATTTTTT 6887

RESULT 4  
US-09-178-973B-8  
; Sequence 8, Application US/09178973B  
; Patent No. 6274710  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFs)  
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543  
; CURRENT APPLICATION NUMBER: US/09/178.973B  
; CURRENT FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 17  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
US-09-178-973B-8

Query Match		14.3%;	Score 686;	DB 3;	Length 7445;
Best Local Similarity		53.8%;	Prod. No. 1.8e-181;		
Matches 2644;		Conservative	0;	Mismatches 1875;	Indels 393;
				Gaps	44;
Qy	29	CTCCCTTCCCAGTCCACCAAGTTGCTCGAGTTAGATTGCTGCANTGCGCGCCCTGCAGAA	88		
Db	2034	CTCTCCCTCTCACTTATCAACTGTTTGACACTTGTGCGATCTCTGATGGCTGTCTCTGCAGAA	2093		
Qy	89	ATCTGTGAGCTCTTCTCTTATATGGGGACCCCTGGCCACACAGCTGCCTCTCTCTCTTGGGCCCT	148		
Db	2094	ATCTATGAGTTTTTCCCTTATATGGGGACTTTGGCCCGCAGCTGCCTGTTCTCATTTGCCCT	2153		
Qy	149	CTTGGTACAGGGAGGAGCAGCTGCGCCCATCAGCTCCCATCTGCAGGCTTGACAAGTCCAA	208		
Db	2154	GTGGGCCACAGGAGCAAAATGCGCTGCCGTCAACACCCGGGTGCAAGCTTGAGGTGTCCAA	2213		
Qy	209	CTTCCAGCAGCCCTATATCACCAACCGCACCTTCATGCTGGCTTAAGAGGTATACATCTC	268		
Db	2214	CTTCCAGCAGCCGTACATCGTCAACCCGACCTTTATGCTGGCCAAAGAGGTACAGCTGCA	2273		
Qy	269	AATCCTGCTCTTCTCGTTGGATCTACTTGGAAATCCAAATAGTTCTTTAAACTTTTCTTCA	328		
Db	2274	TCCTCTTCTCTCCATACCGCCTTGCCATTTTCTCTGAAGCACTTGCNAACCTCTTTAGGG	2333		
Qy	329	GAGCATCTTAAGAGCTTTTAGSAAACCCACTGTTTTATCCCTGAGGCTAGATAAAATTTTCTG	388		
Db	2334	CGCTTTATCTCCGAGGTCTCACTACCTATGTTT-----TCTGTCTCTTTAGAG	2382		
Qy	389	TTTTTTTCAGAGACTCTTTGGGAATCTGGCTTTTTTTTTTTTTTTCTTTGAACTTCTTCTCTCCAT	448		
Db	2383	ACTCTTTAAGGACTGGGTCTTTTTTCTATTCTATTTCGAAGTCTCTCAGGACCATTTTCTCAT	2442		
Qy	449	TTTCGCTTTATGATACATATGATGAATTTTTTCCCAAGAGCGGCCAATCAGTAATCCAT	508		
Db	2443	CTTGGCTTTCAGGACACATATACTGAATTTATCTACAGAGGCGCAATT--AGAAAGCCA	2500		
Qy	509	CTGATGATTTTTTTTTTCTTTATGCCCTCTGTGCAATGTTCTTAAACTCATGCACACATCTG	568		
Db	2501	CCCAGACTGCAATACITTCATTTCTCTGTGCTCTCTTCTGAATCTACTCTCTTGGC	2560		
Qy	569	AATTCTGCTTTTAGTCTTTATGATGTTGCTCTGGGGAGACGGGATGGGGCACAATCTAT	628		
Db	2561	TACTC-----CTGAGACCCACTCGGACATACATCTCTAC	2595		
Qy	629	GTATAAATTTTTTTCTATTGTCTCATGTCCAGACCTTAGTCTTTCTTCTCTCTCCAG	688		
Db	2596	TTACAGGCTTTTCTTCTTCATCTCTTGTGTCACCCAGGCACTTAGGGTTTTC-TCTCTTTTCAG	2654		
Qy	689	GCTAGCTTGCTGATAACAACACAGACGTTCTCTCATTGGGGAGAACTGTTTCCACGGA	748		
Db	2655	GCCAGCCTTGAGATACAAACACAGAGTCCGGCTCATCGGGAGAAACTGTTTCCGAGGA	2714		
Qy	749	GTCAGTGTAAAGCTACAGTTGTGACGAAACAGGGCCGTTGCCGTCATGGGTACTTGGGGT	808		
Db	2715	GTCAGTGTAAAGTCTCACTGTGATGACAGSGC-----TAGCTGGGGAGCT	2761		
Qy	809	GGTGGTCAATGATGTTTTAGTCTTATCCCTTATGACCCCTTTCTGTTTCCCTTCCACTGC	868		
Db	2762	GGTGGACCCCTCTGGGATAG-----TCTGACGATGACCCCTCTGCTTCTTGTCTACCTGC	2817		
Qy	869	AGATGAGTGAGCGCTGCTATCTGATGAAGCAGGTTGCTGAATCTCACCCCTTTGAAGAAGTGC	928		
Db	2818	AGGCTAAGATCAGTGCTACCTGATGAAGCAGAGTGTCTCACTTCACTCCCTGGAAAGCGTTC	2877		
Qy	929	TGTTCCCTCAATCTGATAGGTTCCAGCCTTATATGCAAGGAGGTGGTGCCCTTCTCTGGCCA	988		
Db	2878	TGCTCCCCAGTTCAGACAGGTTCCAGCCCTACATGACAGGAGTGGTACCTTTTCTTGACCA	2937		
Qy	989	GGCTCAGCAACAGGCTAAGCACTATGTAAGTTTCAAGTCTCTCAGGCTATGCCACTTACCC	1048		
Db	2938	AACTCAGCAATCAGCTCAGTCTCTGTGTAAGTCTGACTCTGGGTACCTATGCTCTCTCTCT	2997		

[illegible]

4051	Db	TTTTCTCGCTTCCAGTCCCTTCTACTTTGTAACTTTATTATTGACTGTCTACTATCTG	4111
2121	Qy	GGGAGTTA----CTTTGGTGTCTGTGTAGATATATCTATATATCTAGATGTCAGTT	2176
4111	Db	GTCCATTTACTCGCTTACTGCTGCACCTGTATCTAGCTGGGTCTATAGATTTCTTCAATCTGTG	4170
2177	Qy	TCCAAATCTTTGCCAAATTGTAGAATCTAGAACTGTGTGGGATCTTGTAGCTTGTCTAGTCAAC	2236
4171	Db	TCTAAATTT---GTAAGTCAAAATCTGGAGCTAGCAGAAAGCTTGAAGCTTCCGCACTCTC	4227
2237	Qy	ATAACCTCAGATTTCTGGGATGGTCACTGGCAGAGATAGGGCTAGGAATCAAGGTCTTCCTG	2296
4228	Db	ATGAGCACTTGTCTGGAGGATGGCTTGTGACAGAGTCAATGCTAGAAAGCAGCATCCCTG	4287
2297	Qy	AATCCCAAGCCAGCACTTTTCCGGTGGGTATACAGATTAGTTTTCGGTACCAATTAATTTCT	2356
4288	Db	ATTCCCAAGCTCTGCAC--TTGCCCTAGTGGCCATGTGTAATTTACTTTTGGCTTGATTAAGTAT	4346
2357	Qy	TAGGGAATTTTCAGATTCCTATTGACTCATGTAACTCTGAAAGAGTACTTGTTTTAAAAACA	2416
4347	Db	TTGGGAAA--GCCAGTTCCACGGACCTACATTAATCTGAAGAACCATGCATTGAAACCTA	4404
2417	Qy	GAAAAATGCCCTATGGGCAAAATTTATTTGAAGTCAATTTTGAAGTCAATTAATGCAATTCCT	2476
4405	Db	GAAA---GCTGGGCACAACTTACTAGAGATGATTTTGGAGCTCATTAACCGATGCTC	4460
2477	Qy	TGAACCTTGGAAAGATATAAATCTAGAACAAATGAGAAAGAGCTCGGACTGCATATAGGGCT	2536
4461	Db	TGAAATGTGGCAAAATCAACCCAGATTAACAACAAAGAGCTGGATTTGCAAAATGAGACA	4520
2537	Qy	AATTTCTCGGA-----GTAATAAACACTTATTTTGAAATTAATCAATATCTATCAGATA	2589
4521	Db	AGTATTTTAGAATCACTGGTATTATAGCTATCATCTTAATTAATAATATAGGGCTATATA	4580
2590	Qy	TTGATTATAGTTTAAAAAGCAGAGCAGACAACC--CCGATCTCTTTTATACAGGTTCAAAT	2648
4581	Db	TATATTTAAGATTAAACACAAAGAGTGTAGTCCCTCCCAATTTACTTGGCGTGGTTTCAAA	4640
2649	Qy	AGAGTAAAAATATTAGTAAAGAGATTTATTATAGTTAAATCGAAGTCTGAATTCGTAAGCT	2708
4641	Db	AGAGTAAAAATATCAGTCATGGATTAATATTAGTGTCTCAATGAAATGATGAGATGGAAACCC	4700
2709	Qy	TTTTTTTCTTCTCTCTCCCATCAAGACCTTCCATCTAGTTTCTCTCTTCACTCCCTCA	2768
4701	Db	TTTCCTTACTTTTACCTTCA-----TTTCTTAGTTTTTTTTTTTCTTCAACCTTGA	4752
2769	Qy	ACAAATCCCTAGGAGCATTTATCCATGGTGGCTGGTGATCATTTCTATAGTGAATGAT	2828
4753	Db	TCAAGCCACTAGTAAGCACCTATCTGTGTGAGCTATTATATGACTTTTACAGCAAAACAAC	4812
2829	Qy	ACCATCATGTGGCCCTATTGTGTAAAGRACA--ACAAATGGAAGCTTAGACTAACAAATA	2886
4813	Db	ATTGCTGTGTGGCTCTTTTGGGGAAGGGAACAGATAGCAGGAGGCTCAGCGCTAGCAAGT	4872
2887	Qy	GTGACTCACCCCAAAAACCGAGGAATGATTAGGAGCAGTGAAGTGAAGCTT--GCAAG	2945
4873	Db	CTGACTTCCCTTAAAGCCAGAGGCATGGTTGATAGCAGAGAAAGTGAAGGCTCTTCGCAAG	4932
2946	Qy	CAGGTACAACTAAATACTCAGAAAATGAAGGCTCCAGTTGATGGAAATTTTCAGTAAACAA	3005
4933	Db	TGGGTGTGCTTAAAGTAATCAGAAAACGGAAGGCTCCGCTTGTATGGAATTAATCAGTAAGAT	4992
3006	Qy	GCTTAAACCTTAATTTCCCTCTTTTCCCTCTGTACTTTTAAAAAAGCGTTCTTCCCTGAG	3065
4993	Db	ATCTACCCCTTATCTCTTCTTCTATCGAACCC-----TAAATCGTCTCTTTTCTTGTG	5042
3066	Qy	CATCAATTTAATGAGTGTGACTGTGTTCTTCCCTTGTATAATGAAGGCTTTGTAGTTTAAA	3125
5043	Db	TGTAGGCTGATTAACACACTTGT--TTCTTTTGAAGTTCATGGCTTTGTAGATTTTAA	5100
3126	Qy	TTGTGAAGCCCGATTCTCTGTTATAGAACTATTATCTPAGACATGGAAGGCTGAAATGTTA	3185





QY 1162 ATCATAGAGTATTTGCTTTTGGCTTGTGACTGAGTCACATCTTGAGTTTATAGTGGTGAATG 1221  
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QY 3118 TTCAATTGAGTAATCTTTGAGTTTGTATGAGTGAAGCTTTATTTGTTTATCATGGAA 3177  
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QY 1222 GGGTCTCGAACTTAAAGTGTACAGAAGCCGATTTGGTTTGTCTCGGAAAAAAGCAACTC 1281  
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QY 1282 A-----GGTTGCGTAAGATGAGAAAGGTGTTGG 1309  
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QY 3238 AGAGAAACAGATCTGCTGAGTATAGTACTTATGCGGGGAGCAGGGGCGGATATCCACTGA 3297  
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QY 1310 GAAACATCTAGCTGTGGAATGATCCATTTGAGTCTTAAGTTGTTGAGGGGAGGGGATGG 1369  
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QY 1370 CATGCAGAGAAATTAGAAGAGAAAGTGGGAAATGGGAAGGCTTAAA----- 1415  
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QY 3358 CACTGAGTACNAGTACTTTGGGGGAGGGAATGGCACAGACNAAAGTTGAAGGGAAGG 3417  
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QY 1416 -----GTGCGTGGTGGTTCGGCAGACTGTTGCGCTGTGTGATGTCATGGGA 1460  
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QY 3418 AAGATGAGAGGCTCATGCTGTTGGGGTGTGAAGGTCACTCTTTTCCATGTGATGGAG 3477  
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Db |||||  
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QY 4993 ATCTACCCCTTATCTCTCTCTCTCGAACC-----TAAATCGTCTCTTTTCTTCTG 5042  
QY 3066 CATCATTTAATGAGTGTGACTGTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 3125  
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QY 3126 TTGTGAAGCCAGTTCTTGTGTTATAGAACTATTATCTAGACATGGAGGCTGATGTTA 3185  
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Db |||||  
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QY 3246 GCTTGTGCTTTTAGAAAAAGTGAAGTGTGAGAGAGGAGAAATCTCATGTGTA----- 3296

6206	TGGATACAGTGTGATAATTGA-----	6222
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Qy	4402 TTCATTAGTGTGAGGTGCACCTGAAATTGATGCTGCTGTGGTGGCCT-CTCAGTCCAGAGA	4460
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Qy	4461 GCCGTCAATTTTAAGCTCTTTTGGCAAATCATACAATACTATAAAGGATA-----T	4508
Db	6344 GGCAATGATTTTTAAGCTCTTGGGCAATCATATTATTAFACTCATGCTATAAAATACATTATGTT	6403
Qy	4509 TACTATGAATGTTTTTACAAATGCTTAAAACTCGGTTTTCTGCTCCATCAACCTAAATCTTG	4568
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Qy	4569 CAATTTCT----AATTTGTTCACTTTAGAAAACATGGCATTAANTGCTCAAATACTTTTGCA	4625
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Qy	4626 TTCTTAATTTTCACAGCTTTGAGAGAGATGAGAGATCAAAACAAATGCGAGAACTCGGATTTG	4685
Db	6524 TTCTTAATTTTCATAGCTTTGAGAGAGATGAGAGATCAAGCGAATTTGGGHACTGGAACCTG	6583
Qy	4686 CTGTTTATGTCCTCTGAGAAATGCTGCAATTTGACAGCAAAAGCTGAAAAATGAATAAC	4745
Db	6584 CTGTTTATGTCCTGAGAAATGCTTGGCTCTGAGCGAGAGAGAGCTAGAAAAACGAGAAC	6643
Qy	4746 TAACCCCTCTTCCCTGCTAGAAATACAATTAGATTAGTGCCTCCCAAGCGATTTTTT	4797
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RESULT 6  
US-09-354-243B-8  
; Sequence 8, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; TITLE OF INVENTION: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Ind  
; TITLE OF INVENTION: (TiPs)  
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.1  
; CURRENT APPLICATION NUMBER: US/09/354,243B  
; CURRENT FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:

US-09-354-243B-8

Query Match 14.3%; Score 686; DB 3; Length 7445;  
Best Local Similarity 53.8%; Pred. No. 1.8e-191;  
Matches 2644; Conservative 0; Mismatches 1875; Indels 393; Gaps 44;

Qy	29 CTCCTCCCAGTCCACAGTTGCTCGATTGAGTAATTCCTGCAATGCGCCCTGCAGAA	88
Db	2034 CTCCTCTCTCACATTATCACTGTGTGACACTTGTGCGAATCTCTGATGCTGCTTCGCGA	2093
Qy	89 ATCTGTGAGCTCTTTCTCTTATGGGAGCCCTGGCCACCAGCTGCCTCTCTCTTTGGCCCT	148
Db	2094 ATCTATAGATTTTTCCTTATGGGACTTTTGGCGCCAGCTGCCTGCTTCTCATTTGCCCT	2153
Qy	149 CTTGGTACAGGAGAGAGACGTGCGCCCATCAGCTCCCACTGCAAGCTTGCAAGTCCAA	208



Db 2154 GTGGCCAGGAGCAAAATGCGTCCCGTCAACACCCGGTCAAGCTTGAGGTGTCCAA 2213  
 Qy 209 CTTCCAGAGCCCTATATACCAACCGCACCTTCATGCTGGCTAAGGAGTATACATCTC 268  
 Db 2214 CTTCCAGAGCCGTACATCGTCAACCCGACCTTTATGCTGGCCAAAGGAGTACAGTGC 2273  
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 Db 2274 TCTCTTTCTCTCCAAACCGCCTTCCCAATTTCTCTGAAGCACTTGCAACTCTTTAGGG 2333  
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 Db 2383 ACTCTTTAAGGACTGGGTCTTTTCTATTCTATTCTCAAGGCTCTCAGGACCAATTTCTAT 2442  
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 Qy 1370 CATGGAGAGAAATTAGAAGAGAAAGTGGGAAATGGGAAAGTGGAGGCTTTAA----- 1415  
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 Qy 1881 GAGGTTTCAAAAGTTTCAGCAACATTAAGATAGTTATGCTTGTGTTATTTTATAGCA 1940  
 Db 3874 AGGTATTTGGGCTCCACCGGATAAGATCTGTGTTAGTGA-GTCTGCTTTTATTTTGCAGCA 3932  
 Qy 1941 TATTGAAGGTGATGACCTGATATCCAGAGGAATGTGCAAAAGCTGAAGGACACAGTGAA 2000  
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Db 1679 AATGGCCACAGCAGCAAAAGTTGAAGGGAAAGAGGAGATGAGAGGCGCTCAATGTGGGG 1738  
Qy 1451 TGTCAATGGGAAGCCACAAAATCGAGGCGGTGTGAATTTGATCGCGCTGAACATTTGAAC 1510  
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Db 1918 GAGTTAAGCTCAGTTTCTTCT-----ACCGTCTTGTCTAC 1956  
Qy 1691 AAATGCAAACTCAGTAGGATTTCCAAAGATGAAGAGAGGTCTCTTTGTAAGGGAAGTGA 1750  
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Qy 1871 CATGGGTGTGGAGGTTTCATAAAGTTTCAGACAACTAATTAAGATAGTTATCTTTATTG 1930  
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Db 2551 AGCATCCCTGATTTCCAGCTCTGCAC-TTGCTAGTGGCCACGTTGATTTACTTTAGCCT 2609  
Qy 2347 CATTAAATCTTAGGAAATTCAGATTTCTTATTTGACCTCAGTCAATCTGTAAGTACTTGA 2406  
Db 2610 GATTAAATTTTGGGAAA--GCCAAATTTCCACCGACCTCATATTCGGAAGAGCATGCA 2667  
Qy 2407 TTTAAAAACAGAAAATGCTTATGGGCAATTTATTTGAAGTCAATTTTGAAGTCATTAA 2466  
Db 2668 TTGAAAACACTAGAA-----GCTGGGCACAAATCTACTAGATGATTTTGGAGCTCATTA 2723  
Qy 2467 TGCATTTGTAACCTTGAAGAAATAAATCTAGAACAAATGAGAAAAAGAGCTGGACTGC 2526  
Db 2724 ACTGATGCTCTGAAATGTGATCAATCAACCCAGAAATAACAAACAAAGAGCTGGATTGC 2783

Qy 2527 ATATAGGCTTAATTTCTGGAGTAATAAACACATTAT-----TTTGAATTTATCATAATA 2578  
Db 2784 AATAGGACAAAGTATTTTAGAATCACTGGTATTAAACAGCTGTCTATTAATTTAAATATAG 2843  
Qy 2579 TCT---ATCAGATATTGATTTAGTTTAAAGCAGAGCAGACAAC-CCCGATCTCTTTT 2634  
Db 2844 TGTCTATTTAGCTGCCCTATTTAAGATTTAAACACAAAGAGTGGATAACTTCCCAATTTACTG 2903  
Qy 2635 ATACAGGTTCAATAGAGTAAATAATTTAGTAAGAGATTTATTATAGTTAAATGGAAGTC 2694  
Db 2904 GGCTGTTTTCATAGAGTAAATAATCAGTCATAGATTTATTTATAGTGTCAATGAAGTA 2963  
Qy 2695 TGAATTTGGTAAGCTTTTTTTCTTCTCTCTCCCATCAAGACCTTCCATTTCTAGTTCTT 2754  
Db 2964 TGAGTTGGAAACC--CTTTCTTCTTCTTTTACCTTCTTCTTCTTCTTCTTCTTCTTCT 3020  
Qy 2755 CTTTCTCTCTCAAAATCTCCTAGGAGCATTTATCCATGTTGGCTGGCTGGTGTACATTT 2814  
Db 3021 TCTTCAACCTCTGATCAAGCCTAGTAAGCCTATCTGCTGCGAGCTATTTATATGACT 3080  
Qy 2815 CTATAGTGAATGATACCATCATGTGGCTTATTTGGTGAAGAAACA--ACAATGGAAGGC 2872  
Db 3081 TTACAGCAACAACATTTGCTGTGTGGCTCTTTTGGGAAGGGAACAGGATAGCAGAGGC 3140  
Qy 2873 TTAGACTTAACA--TAGTGACTCAACCCAAACCGGAGGAATGATTAGGAGCAGTGAAAGT 2931  
Db 3141 TCAGGCTAGCAAGTCTGGACTCAACCTAAAGCCAGAGGCATGTTGATAGCAGAGAAAGT 3200  
Qy 2932 GAGCTCTTT-GAAGCAGGTACAACCTAAATACTCAGAAAACATGAAGGCTCAGTTGATGG 2990  
Db 3201 GAGGCTCTTCAACAAGTGGGTGCTTTAAGTAATCAGAAAACAGGAAGGCTCTGGTTGATGG 3260  
Qy 2991 AATTTTCAGTAACAAGCTTAACCTTAATTTCCCTTTTCCCTCTTGACTTTTTAAAAA 3050  
Db 3261 AATTATCAGTAAGATTTACTACCTTTATCTCC-----TTCTTCTATAGAGCTTAACCG 3313  
Qy 3051 GCCTTTCTTCTCCTGAGCATCATTTAATGAGTGTGACTGTTTCTTCTTCTTCTTCTTCTTCTTCT 3110  
Db 3314 TCTCTCTTCTTCTGTGTAGGCTGATAACACGCTTGT--TTCTTTTGTAGTTTCATGG 3371  
Qy 3111 CTTTGTAGTTTAAATTTGAAGCCAGCCAGTCTCTGTTTATAGAACTATTTATCTAGACATG 3170  
Db 3372 CTTTGCAGATTTTTCAGTGTCTGCGAGTCTTCTGT--TAGAGGTTTGTGTACCTTTGACCC 3429  
Qy 3171 GAGGCTGATTTAGCATGCCACAGAGGCATGCTTTTACACATCTTCTGAAATGCTGTGTAAGAAAT 3230  
Db 3430 TGGGCTTGGATTTAGCATGCCAAAGGCACACACTTCTGAAATGCTGTGTAAGAAAGTTAT 3489  
Qy 3231 TACTGATTTCTATCTTGTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 3290  
Db 3490 TATTCATTTACT-----TTGCTTTTGGAAAGGTGAAGTGTGTGTGAGAAAGAACTCA 3541

## RESULT 8

US-09-419-568F-29  
: Sequence 29, Application US/09419568F  
: Patent No. 6331613  
: GENERAL INFORMATION:  
: APPLICANT: Dumoutier, Laure  
: APPLICANT: Louhed, Jamila  
: APPLICANT: Renaud, Jean-Christophe  
: TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
: TITLE OF INVENTION: (TIPs) The Proteins Encoded, and Uses Thereof  
: FILE REFERENCE: LUD 5543.2  
: CURRENT APPLICATION NUMBER: US/09/419,568F  
: PRIOR FILING DATE: 1999-10-18  
: PRIOR FILING DATE: US09/354,243  
: PRIOR FILING DATE: 1999-07-16  
: PRIOR FILING DATE: US09/178,973  
: NUMBER OF SEQ ID NOS: 29  
: SEQ ID NO 29

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; LENGTH: 5935
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-09-419-568P-29

Query Match      13.6%; Score 650; DB 3; Length 5935;
Best Local Similarity 56.5%; Pred. No. 2.1e-171;
Matches 1863; Conservative 0; Mismatches 1285; Indels 152; Gaps 29;

Qy 29 CTCCTCCCGCAGTCACACAGTTGCTCGAGTTGCTGAGATTGCTGCAATGGCGCCCTGCGAGAA 88
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 356 CTCCTCCCTCAGTTATCAACTTTTGACACTTTGTGCGATCGGTGCGTGTGCTCTGTCGACAA 415
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 89 ATCTGTGAGCTCTTTCTTATGGGAGCCCTGGCCACACAGCTGCCCTCTCTCTTGGGCCCT 148
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 416 ATCTATGAGTTTTCCTTATGGGAGCTTTGGCGCCAGCTGCTCTCTCTCAATGGCCCT 475
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 149 CTTGGTACAGGGAGGAGCAGCTGGCGCCCATCAGCTCCCACTGCAGGCTTGCAAGTCCAA 208
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 476 GTGGGCCCGCAGGAGCAATGGCTGCCCATCAAACCCGGGTGCAAGCTTGAGGTGTCCAA 535
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 209 CTTCCAGCAGCCCTATATACAAACCGCACTTCATGCTGGCTAAGGAGGTATACATCTC 268
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 536 CTTCCAGCAGCCGTACATCGTCAACCGCACCTTTATGCTGGCCAGGAGGTACAGCTGCA 595
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 269 AATCTGCTCTTTCTCGTTGGATCTACTTGGAAATCCAAATAGTTCTTAAATTTCTTCA 328
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 596 TCTCTTTCTCCATACCGCCTTGGCAATTTCTCTGAAAGCACTTGCAAACTCTTTAGGGGC 655
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 329 GAGCATCTTAAGAGCTTTAGGAAACCCACTGTTTATCCCTGAGGTAGATAAATTTCTG 388
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 656 GCTTATCTCCGAGGCTCTACTACCTATGTTTCTGTCT-----CTTTAGAG 703
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 389 TTTTTCAGAGCTCTTTGGGAAATCGGCTTTTTTTTTTTTCTGAACTCTTCTCTCCAT 448
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 704 ACTCTTTAAGCACTGGATCTTTTCTTCTATTTCTAATTTCAAGGCTCTCAGGACCATTTCTAT 763
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 449 TTTGGCTTTATGATACATATGATGAAATTTTCCCAAGAGCGCCATTCAGTATCCAT 508
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 764 CTTGGCTTTCAGGACACATATAGTAATTTATCTACAGAGCGCGT---AGAAAGCCA 821
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 509 CTGATGATTTTTCCTTATGCTCTGTGCAATGTTCTTAAACTCATGACACATCTG 568
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 822 CCCAGACTGCAATCTTCCATCTGTGTGCTCTTCTGAACTCATATCTCTTGGC 881
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 569 AATCTGCTTTTAGTCTTTATGATGTGCTCTGGGAGACGGGATGGGACATGTCTAT 628
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 882 TACTC-----CTGAGACCCACTGGCGGACATACATCTCTAC 916
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 629 GTATAAATTTTTCATTTGCTCAATGTCAGACCCCTTAGTCTTTTCTCTCTTCCAG 688
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 917 TTACAGGCTTTTCTTCCATCTCTTGTCAACCCAGCACTTAGGGTTTC-TCTCTTTCAG 975
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 689 GCTAGCTTGGCTGATAACAAACAGACGCTCGTCTCATTTGGGAGAACTGTTCACGGA 748
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 976 GCCAGCTTGAGATAACAAACAGACGCTCGGCTCATCGGGAGAACTGTTCGAGGA 1035
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 749 GTCAGTGAAGCTACAGTTGTGAGAAACAGGGCGGTGTGCGCTCCATGGGTACTTGGGGT 808
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1036 GTCAGTGAAGTCCCTCACTGTGATGAGCAGGCG-----TAGCTGGCGGAGCT 1082
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 809 GGTGGTGAATGGTTTAGTCTTATCCCTTATGACCCCTTTCTGTTTCCCTTCCACCTGC 868
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1083 GGTGGACCCCTCTGGGATAG-----TCTGACGTATGACCCCTGCTGCTTCTGTACCTGC 1138
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 869 AGATGAGTGAGCGCTGCTATCTGATGAAGCAGGTGCTCAATTCACCCCTTGAAGAGTGC 928
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1139 AGGCTAAGCATCAGTGTCTCTCTGATGAAGCAGGTGCTCAATTCACCCCTGGAAGCATTC 1198
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 929 TGTTCCTTCAATCTGATAGTTTCCAGCCTTATATGACAGAGGTGGTCCCTTCTTGGCCA 988
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1199 TGCTCCCCCAGTCAGACAGGTTCCGGCCCTATATGACAGAGGTGGTGGCTTTCTTCTGACCA 1258
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

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Qy 989 GGCTCAGCAACAGGCTAAGCACAATGTGTAGTTGAGTTCAGCTCTCAGCCTATGCCCCACCTACCC 1048
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1259 AACTCAGCAATCAGCTCAGCTCCTGCTGTGTAGTCTGGCTCTGGCTACCTATGCTCTCTCT 1318
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1049 CTCCTTCCCTCTTCCACAGAGACCCCTTACCACCACTCTCTCTCTCTCTCTCTCTCTCTCTCT 1108
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1319 CTTCTCTCTCTTATTTCCAGTAAGAACCCGAGGTCTGCTCCCTCTCTCTCTCTCTCTCTCTCTCA 1378
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1109 TAGCTAGCAGGAGAGAGTGTCTTGGCAGCAGTGTATCAGGAGTCA-----TTTGGG 1161
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1379 GAGGGGCTCAGCACCAACCACCATCATAGGCCACTTGAAATAGGTACAAAGGCTTTGGC 1438
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1162 ATCATAGAGTATTTGCTTTTGTCTTGTGACTCAGTCACATCTTGTAGTTTATAGTGGTGAATG 1221
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1439 TTTCAATTGAGTAATACCTTTGAGTTTGTATTAAGTTTATTTGTTTATCCATCGAA 1498
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1222 GGCTGTGAACTTAAAGTGTACAGACCGCATTTGGTTTGTCTTCGGGAAAAAAGCAACTC 1281
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1499 AGAAATCAACTCAAAATTTCTGTAGGATGAGAAAGATGTTGGGAACGAAAAAAGGCTAGAT 1558
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1282 AGTTTGGTAA---GATGAGAAAGTGTTCGGGAAACATCTAGCTGTGGAAATGGAATCCA 1338
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
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Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
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Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1399 AAATGGGAAGCGCTTAAAGTCGGTGGTGGTGGCGGAGACTGTTGCC-----TGTGTA 1450
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1679 AATGGCACAGAGCAAAAGTTGAAGGGGAAAGAGGAGATGGAGAGGCCCTCAATGTTGGGG 1738
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1451 TGTTCATGGGAAGCCCAAAATCGGAGCGGTGAACTTGTGATCGCGCTGAAACATTTGAAAC 1510
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1739 TGTGAAAGGTCACCTCTTTTCCATGTGATGGAGGTTAAGAAAAATCAGTGTGTAGTT 1798
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1511 TATGAAAAAAGTTTGTAGTGGAGTGGGCCCAAGTAAAAAGGCCCTAGGACTTACTGAAGAGG 1570
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
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Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1571 GCTTAATTTTCAATGAGATGTTTTATGTACATTTCTTGTCTTAAGCATGCAATTTCTG 1630
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1858 CGCGGCTTTTCAACGAGAACTTTATGCTCATCTCTTGTGTACACTCCACACTTTGAT 1917
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1631 GAGATAGATGAGGTTTATTTCTTACAGAAATTTGCATAACTACTCTCCGCTCTTTCCAC 1690
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1918 GAGTTAAGCTCAGGTTTCTTCT-----ACCGTTCCTTGTGTAC 1956
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1691 AATGCAAACTCAGTAGGATTTCCCAAAGATGAAGAGAGTCTCTTGTAGGGGAAGTGA 1750
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1957 TGGTGGAACTTCAGTAGGATTTCCCAAAGACGAGGACAGCTCTTCTGTAGGGGAGGAC 2016
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1751 CTGGAATCTGCGCTCAAGGGAATTCAGAGCTCAGGAAATCTAGGTCACTGTTTGAATC 1810
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 2017 CTGGAATTCAGTGTCTTAGAGAACGAAATAGCTCAGAGAAATCTAGGTCAACGCTGAAATCT 2076
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1811 TAGTCTAATGTGGGCAAAATTAATAAGAGCTTTTAATTTCCAGGTGAATTTGACTTACTCTC 1870
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 2077 AGTTCACAGCGGGCAAAATTAAGTGAACGCTCTTAATTCAGGTGAACGGTCACGTGCTC 2136
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 1871 CATGGTGTGAGGTTTCATAAAGTTTCAGCACAACTTAAGATAGTTATGCTTGTATTG 1930
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 2137 AGATATACTGAGGTATTGGGCTCCACCGGATAGATTTCTGTAGTGA-GTCTGCTTTTA 2195
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Qy 1931 TTTTATAGCATATTGAAGGTGATGACCTGCAATATCCAGAGGAATGTGCAAAAGCTGAAGG 1990
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 2196 TTTTGCAGCATCAGTGGTGAACGACCAAGAACATCCAGAGAAATGTCAAGAGGCTGAAGG 2255
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Qy 1991 ACACAGTGAAGGAGGTAGGACTGATACTGTCAATGTCTAGTCTATGCAATAGGAGAGACA 2050
Db      ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Qy 2256 AGACAGTGAAGGAGGTACTATTGGCAAGGCCAATACTAAGCCATTCACTAGTAGGAGACGTG 2315
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QY 2051 AATGTTGTTTCTTCTTCTTCTTCCATCACATTTGTGATTTTTTCACTTGATCTC 2110
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QY 2374 CTACTGTCTGTCATTTACTCATTCTAGCTGACCTGCACTCTAGCTGGGTCTATAGATCTT 2433
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QY 2167 GATGTCAGTTTCCAAATCTTTGCAAATTTGTAGAAATTTCTAGAACTGGTTGGGATCTTAGCTTT 2226
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QY 2434 TCAATCTGTGCTAAATTT---GTAAGTCAAAATCTTGGAGCTAGCAGAAAGCTTAGCTC 2490
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QY 2227 GTCTAGTCAATACCTCAGATTTCTGGGGATGFTCACTGGCAGAGATAGGCTTAGAATGC 2286
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QY 2491 AGCCAGTCTCATGAGCACTTTGCTCGAGATGGCTTTGTGACAGAGTCAATGCTTAGAAGAC 2550
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QY 2287 AGGTCTCTGAATCCCAAGCCAGCACTTTCCCGTGTGTATACAGATTAGTTTGTGTAC 2346
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QY 2551 AGCATCCCTGATCCCAAGCTTGAC--TTGCTAGTGGCCACGTTGTAATTTACTTTAGCCT 2609
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QY 2724 ACTGATGCTCTGAATGTGATCAATCAACCCAGATTAACAACAAAGAGCTGGATTTGC 2783
Db      |||||
QY 2527 ATATAGGCTCAATTTCTGGAGTATAAACAATTAT-----TTTGAATTTATCATATA 2578
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QY 2784 AATAGACAAAGTATTTAGATCACTGGTATTACAGCTGTCTATTTAATTAATAATAG 2843
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QY 2579 TCT---ATCAGATATGATTATAGTTTAAAGCAAGAGCAGACAAC--CCGATCTCTTTT 2634
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QY 2844 TGTCTATTAGTCTGCTTATTTAAGATTTAAACACAAAGAGTGGATAACTTCCCAATTTACTG 2903
Db      |||||
QY 2635 ATACAGTTCATAATAGATTAATAATATTAGTAGAGATTTTATATAGTTAAATGGAGTC 2694
Db      |||||
QY 2904 GGCCTGTTTCAATAGAGTAAATAATATCAGTCATAGATTAATATATAGTGTGATGAAGA 2963
Db      |||||
QY 2695 TGAATTTGGTAAAGCTTTTCTTCTCTCTCTCCATCAAGACCTTCCATTTCTAGTTCTTT 2754
Db      |||||
QY 2964 TGAGTTGGAACC---CTTTCTTACTTTTACCTTCATTTCTTAGTTATTTATTTTTTTT 3020
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Db      |||||
QY 3201 GAGGCTCTTCAACAAGTGGGTGTGCTTAAGTAATCAGAAAACAGGAAGGCTCTGGTTGATGG 3260
Db      |||||
QY 2991 AATTTTCAGTAAACAAGCTTAACCTTAATTTCCCTTTTCTTCTTCTTGAACCTTTCTTCCAT 3050
Db      |||||
QY 3261 AATTTATCAGTAAGATATCTACCTCTTATCTCC-----TTCTTCTATAGAAGCTTAACCG 3313
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QY 3051 GCGTTTCTTCTGAGCATCATTTAATGAGTGTGACTGTCTTCTCTTCTTGTATTAATGAAGG 3110
Db      |||||
QY 3314 TCTCTCTCTTCTGTGTAGGCTGATTAACAAGCTTGT---TTCTTTTGTAGTGTCTATGG 3371
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QY 3111 CTTTGTAGTTTAAATTTGTGAAGCCAGTTTCTCTTGTATTAGAACTATTATCTAGACATG 3170
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## RESULT 9

US-09-354-243B-29

; Sequence 29, Application US/09354243B

; Patent No. 6359117

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Louheid, Jamila

; APPLICANT: Renauld, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa

; TITLE OF INVENTION: (TIFF)

; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5543.1

; CURRENT APPLICATION NUMBER: US/09/354,243B

; CURRENT FILING DATE: 1999-07-16

; PRIOR APPLICATION NUMBER: US09/178,973

; PRIOR FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 29

; SEQ ID NO 29

; LENGTH: 5935

; TYPE: DNA

; ORGANISM: Homo sapiens

; FEATURE:

US-09-354-243B-29

Query Match

Best Local Similarity 13.8%; Score 650; DB 3; Length 5935;

Matches 1863; Conservative 0; Mismatches 1285; Indels 152; Gaps 29;

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QY 29 CTCCTTCCCAGTCCACAGTTGCTCGAGTTAGAAATGCTGCAATGCGCCGCTGCAGAA 88
Db      |||||
QY 89 ATCTGTAGTCTTCTCTTATGGGACCCCTGGCCACAGCTGCTCTCTCTTGGCCCT 148
Db      |||||
QY 416 ATCTATAGTCTTCTCTTATGGGACTTTGGCGCAGCTGCTGCTCTCTCATGCGCT 475
Db      |||||
QY 149 CTTGGTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGAGGCTTGACAAAGTCAA 208
Db      |||||
QY 476 GTGGGCCAGGAGGCAAAATGCGCTGCCCATCAACACCGGTGCAAGCTTGAGGTGTCAA 535
Db      |||||
QY 209 CTTCCAGCAGCCCTATATACCAACCGCAGCTTTCATGCTGGCTAAGGAGGTATACATCTC 268
Db      |||||
QY 536 CTTCCAGCAGCCGCTATCATGCTCAACCGCACCTTTATGCTGGCCAGGAGGTACAGCTGCA 595
Db      |||||
QY 269 AATCTGCTCTTCTCTGTTGGATCTACTTGGAAATCAAAATAGTTCTTAAACTTTCTTCA 328
Db      |||||
QY 596 TCTCTTCTCTCCATACCGCTTGGCAATTTCTCTGAAGCACTTGCAAACTCTTTAGGGGC 655
Db      |||||
QY 329 GAGCATCTCTAAGAGCTTTTAGGAACCCACAGCTGTTTATCCCTGAGGGTAGATAAATTTCTG 388
Db      |||||
QY 656 GCTTTATCTCCGAGGCTCTCACTACCTATGTTTTTCTGTCT-----CTTTAGAG 703
Db      |||||
QY 389 TTTTTCAGAGACTCTTTGGGAATCTGGCTTTTTTTTTTTTTTTTCTTGAACCTTCTCTTCCAT 448
Db      |||||
QY 704 ACTCTTTAAGGACTGGAATCTTTTCTTATTTCTATTTTCAAGGTCTCAGGACCATTTCTCTAT 763
Db      |||||
QY 449 TTTGGCCTTTATCATATATATGATTAATTTTCCCAAGAGCGGCCATTCAGTAATCCAT 508
Db      |||||
QY 764 CTTGGCCTTCAGAGACATATATGTAATTTTATCTACAGAGGCGCGTTT--AGAAAGCCA 821
Db      |||||
QY 509 CTGATGATTTTTTTTTTCTTTTATGCTGCTGTCATTTGTTCTTAAACTCATGACACATCTG 568
Db      |||||
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[illegible]

Db	1958	CGGGCTTTTCACAGAGAACTTTATGCTCATCTCTTGTGCTACACTCCACCTTGTAT	1911
Qy	1631	GAGATACGATTGAGGTTTTATTCTTTCACAGAAATTTGCATAAACTACTCCGCTCTTTCCAC	1690
Db	1918	GGGTTAAAGCTCAGGTTTCGTTTCT-----ACCGTCTTGCTAC	1956
Qy	1691	AAATGCNAACCTCAGTAGGATTTCCAAAGATGAAGAGAGGTCTCTTGTAAAGGAAGTGA	1750
Db	1957	TGGTGGAAACTTCAGTAGGATTTCCCAAGACAGGACAGCTCTCTCTGTAAAGGAGGAGAC	2016
Qy	1751	CTGGATTCTCGCTCCAAAGGGAATTCAGAGGCTCAGAAATCTAGGTCACTGTTGGAATC	1810
Db	2017	CTGGAATTCAGTGTCTTAGAGAACGAATAGCTCAGAGAACTAGGTCAACGTGGAATCT	2076
Qy	1811	TAGTCAATTTGTGGGCAAAATTACTAAGAGCTTTAAATTCAGGTGAATTTGACTGTACCTC	1870
Db	2077	AGGTCAAGCGGCAAAATGACTGAACGCTCTATTCCAGGTGAACGGTCAAGTGCCTC	2136
Qy	1871	CATGGGTGCGAGGTTCAATAAGTTTCAGCAAAACATTAAAGATAGTTATGCTTGTATTG	1930
Db	2137	AGATATACTGAGGTATTGGGCTCCCAAGGATAGGATTTCTGTTAGTGA-GTCTGCTTTTA	2195
Qy	1931	TTTTATAGCATATTGAAGTGATGACTCGCATATCCAGAGGAATGTCAAAAGCTCAAGG	1990
Db	2196	TTTTGCAGCACATCAGTGGTGAAGCAGCAAGAACATCCAGAAAGATGTCAGAGGCTGAAG	2255
Qy	1991	ACAGGTGAAAAAGGTAGGACTGATAACTGCTCAATGCTAAAGTCAATGCAATAGGAGAGACA	2050
Db	2256	AGACAGTGAAGAGGTACTATTTCGCAAGCCACAACTAAAGCCATTTCAGTAGGAGCGTG	2315
Qy	2051	AATGTTGTTTTCTTCCTTTCTTCTTCCATCACTTTGTGATTTTTTCATTGATCTC	2110
Db	2316	GGGATTTCTTCTCTGCTTCCAGTCT--CTTCTACTTTGTAACTTTCTTTGACTTGT	2373
Qy	2111	CTACCACAGGCGATT----ACTTTGGTCTGCTGTAAGTCTAGATATATCTATATCTTA	2166
Db	2374	CTACTGTCTGTCATTTACTTACTCTTAGCTAGCTGCACCTGCATCTAGCTGGGCTATAGATCTT	2433
Qy	2167	GATGTCAGTTTTCAAATCTTGCAAAATTTGTAGAATTTCTAGAACTGGTTGGGATCTTAGCTT	2226
Db	2434	TCAATCTGTCTTAATTT---GTAAGTCACAATTTCTGGAGCTAGCAGAAAGCTTAGCTC	2490
Qy	2227	GTCPTAGTCACATAACCTCAGATTTCTGGGATGTCTAGTGGCAGAGATAGGGCTAGAAATGC	2286
Db	2491	AGCCAGTCTCATGACACTTGTCTCGGAGGATGGCTGTGACAGAGTCAATGCTTAGAAGAC	2550
Qy	2287	AGGTCTCTGNAATCCCAAGCCAGCACTTTTCCCGGTGGTGATACAGATTAGTTTTCGTAC	2346
Db	2551	AGCATCCCTGATTTCCAGTCTCTGCAC--TTGCCCTAGTGGCCACGTGAAATTACTTTAGCCT	2609
Qy	2347	CATTAAATCTTTAGGAAATTTTCAGATTCTCTATTGACTCATGTAATCTGAAGAAGTACTTG	2406
Db	2610	GATTTAGTATTGGGAAA--GCCAATTCACCCGACCTACATTAATCCGAGAGAGCATGCA	2667
Qy	2407	TTTAAAAACAGAAAAATTCGCTATGGGCCAAATTTATTGGAAGTCATTTTGAAGTCATTA	2466
Db	2668	TTGAAAACCTAGAAA---GCTGGGCACAAACTTACTAGAGATGATTTTTTGGAGTCAATTA	2723
Qy	2467	TGCATTCTGTTGAAACTTTGGAAGAAATTAACCTCAGAACAAATGAGAAAAGAGCTGGACTGC	2526
Db	2724	ACTGATGCTCTGAAATGTGTATCAAACTCAACCCAGAAATTAACAAAAGAGAGCTGGATTTGC	2783
Qy	2527	ATATAGGCTAAATTTCTGGAGTAAATAAACACTTAT-----TTTGAATTAATCAATAA	2578
Db	2784	AAATAGGACAAAGTATTTAGAAATCACTGGTATTAAACAGCTGTCTATTTAATTAATAATAG	2843
Qy	2579	TCT---ATCAGATATTGATTATATAGTTTAAAGCAAGAGCAGACAAC--CCGATCTCTTTT	2634
Db	2844	TGTCATTTAGCTGCCCTATTTAAGATTAAACACAAGAGTGGATAAATCTCCCAATTTACTCG	2903
Qy	2635	ATACAGGTTCAATAGAGTAAAAAATATTAGTAAAGATTTTATTATAGTAAATGGAAGTC	2694
Db	2904	GGCCTGGTTTTCAATAGAGTAAAAAATATCAGTCAATAGATTAATTAATATAGTGCATGAAGA	2963







Db 601 TGTGTGATTTTCAAGACCTTTTAATCCATTTTGAAGAATCAATTTTATATTTGCAATGGG 542  
Qy 3359 TTGCCATGTGGAAGAGTATATGCTTTTGTCTGGTAGCTTTCAGAAAGCACAGGAGGGA 3418  
Db 541 TTGCCATGTGGAAGAGTATATGCTTTTGTCTGGTAGCTTTCAGAAAGCACAGGAGGGA 482  
Qy 3419 GAGCAATGTCTTCAGAGAAAGATCAACAGGAGGAGAAACTGTGAGAGCTGTCTGAAATA 3478  
Db 481 GAGCAATGTCTTCAGAGAAAGATCAACAGGAGGAGAAACTGTGAGAGCTGTCTGAAATA 422  
Qy 3479 GGGTGGTTTTGGGAGGCATTAATTTCCCTCTCGTTTGGGGGTAAGAGCGCAGGTTGG 3538  
Db 421 GGGTGGTTTTGGGAGGCATTAATTTCCCTCTCGTTTGGGGGTAAGAGCGCAGGTTGG 362  
Qy 3539 TAGTAAATGTCATGACAGACAGTAGTAGGGGACGATAAACTTTAAATTTCTTTATAGTCTTGG 3598  
Db 361 TAGTAAATGTCATGACAGACAGTAGTAGGGGACGATAAACTTTAAATTTCTTTATAGTCTTGG 302  
Qy 3599 AGTCTTTGAGATAGAAAGATATCTTTTGGCCCTTATGTCAAAAGAAAGTATGAAAGGT 3658  
Db 301 RGTCTTTGAGATAGAAAGATATCTTTTGGCCCTTATGTCAAAAGAAAGTATGAAAGGT 242  
Qy 3659 GAAAGGGCGGAAGAAAGCAGGAAAGGAAGCAACCATGTATTATATAGAGCAATGGTGA 3718  
Db 241 GAAAGGGCGGAAGAAAGCAGGAAAGGAAGCAACCATGTATTATAGAGCAATGGTGA 182  
Qy 3719 CAAGGTTTTCTGAAATAATGCAAAATATGATATAGATAGAGAAATTCAGTAGGGAATGC 3778  
Db 181 CAAGGTTTTCTGAAATAATGCAAAATATGATATAGATAGAGAAATTCAGTAGGGAATGC 122  
Qy 3779 TTTTCACATTGAATTTGGGTTTCCCTTCGATTAAGTTTGGATCCTCATCTGCATTGAC 3838  
Db 121 TTTTCACATTGAATTTGGGTTTCCCTTCGATTAAGTTTGGATCCTCATCTGCATTGAC 62  
Qy 3839 TTGAGAGAGAAAGAAATGAATGTTAGGACCTATATCTGTTTTTCTATTAACTAAAGCAAG 3898  
Db 61 TTGAGAGAGAAAGAAATGAATGTTAGGACCTATATCTGTTTTTCTATTAACTAAAGCAAG 2  
Qy 3899 T 3899  
Db 1 T 1

RESULT 12  
US-09-419-568F-24  
; Sequence 24, Application US/09419568F  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (Tips) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543-2  
; CURRENT APPLICATION NUMBER: US/09/419,568F  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 24  
; LENGTH: 690  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-09-419-568F-24

Query Match 5.4%; Score 258; DB 3; Length 690;  
Best Local Similarity 100.0%; Pred. No. 9e-62;  
Matches 258; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TGCAACAGCAGAAATCTTTGAGAACAGGTTCTCTTCCCTCCAGTCACCAAGTTCTCGAGTTAG 60  
Db 1 TGCAACAGCAGAAATCTTTGAGAACAGGTTCTCTTCCCTCCAGTCACCAAGTTCTCGAGTTAG 60  
Qy 61 AATTGTCTGAATGGCGCCCTCGAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGG 120  
Db 61 AATTGTCTGAATGGCGCCCTCGAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGG 120  
Qy 121 CCACCAAGTCGCTTCTCTTGTGCGCCCTCTTGTGTACAGGAGGAGCAGCTGCGCCCATCA 180  
Db 121 CCACCAAGTCGCTTCTCTTGTGCGCCCTCTTGTGTACAGGAGGAGCAGCTGCGCCCATCA 180  
Qy 181 GCTCCCACTGCAGGCTTTGACAAAGTCCAACTTCCAGCAGCCCTATATCAACCAACCGCACCT 240  
Db 181 GCTCCCACTGCAGGCTTTGACAAAGTCCAACTTCCAGCAGCCCTATATCAACCAACCGCACCT 240  
Qy 241 TCATGCTGGCTAAGGAGG 258  
Db 241 TCATGCTGGCTAAGGAGG 258

RESULT 13  
US-09-354-243B-24  
; Sequence 24, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa  
; TITLE OF INVENTION: (Tips)  
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.1  
; CURRENT APPLICATION NUMBER: US/09/354,243B  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 24  
; LENGTH: 690  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-09-354-243B-24

Query Match 5.4%; Score 258; DB 3; Length 690;  
Best Local Similarity 100.0%; Pred. No. 9e-62;  
Matches 258; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TGCAACAGCAGAAATCTTTGAGAACAGGTTCTCTTCCCTCCAGTCACCAAGTTCTCGAGTTAG 60  
Db 1 TGCAACAGCAGAAATCTTTGAGAACAGGTTCTCTTCCCTCCAGTCACCAAGTTCTCGAGTTAG 60  
Qy 61 AATTGTCTGAATGGCGCCCTCGAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGG 120  
Db 61 AATTGTCTGAATGGCGCCCTCGAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGG 120  
Qy 121 CCACCAAGTCGCTTCTCTTGTGCGCCCTCTTGTGTACAGGAGGAGCAGCTGCGCCCATCA 180  
Db 121 CCACCAAGTCGCTTCTCTTGTGCGCCCTCTTGTGTACAGGAGGAGCAGCTGCGCCCATCA 180  
Qy 181 GCTCCCACTGCAGGCTTTGACAAAGTCCAACTTCCAGCAGCCCTATATCAACCAACCGCACCT 240  
Db 181 GCTCCCACTGCAGGCTTTGACAAAGTCCAACTTCCAGCAGCCCTATATCAACCAACCGCACCT 240  
Qy 241 TCATGCTGGCTAAGGAGG 258  
Db 241 TCATGCTGGCTAAGGAGG 258

RESULT 14  
US-09-949-016-5443  
; Sequence 5443, Application US/09949016

; Patent No. 6812339  
; GENERAL INFORMATION:  
; APPLICANT: VENTER, J. Craig et al.  
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED  
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF  
; FILE REFERENCE: CL001307  
; CURRENT APPLICATION NUMBER: US/09/949,016  
; CURRENT FILING DATE: 2000-04-14  
; PRIOR APPLICATION NUMBER: 60/241,755  
; PRIOR FILING DATE: 2000-10-20  
; PRIOR APPLICATION NUMBER: 60/237,768  
; PRIOR FILING DATE: 2000-10-03  
; PRIOR APPLICATION NUMBER: 60/231,498  
; PRIOR FILING DATE: 2000-09-08  
; NUMBER OF SEQ ID NOS: 207012  
; SOFTWARE: Fast-Seq for Windows Version 4.0  
; SEQ ID NO 5443  
; LENGTH: 689  
; TYPE: DNA  
; ORGANISM: Human  
; US-09-949-016-5443

Query Match 5.3%; Score 256; DB 3; Length 689;  
Best Local Similarity 100.0%; Pred. No. 3.3e-61;  
Matches 256; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 3 CACAAGCAGAAATCTTCAGAACAGAGTTCTCTCCCTCCAGTCACCAAGTTCGAGTTAGAA 62  
Db 2 CACAAGCAGAAATCTTCAGAACAGAGTTCTCTCCCTCCAGTCACCAAGTTCGAGTTAGAA 61  
Qy 63 TTGTCTGCAATGGCGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGAGCCCTGGCC 122  
Db 62 TTGTCTGCAATGGCGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGAGCCCTGGCC 121  
Qy 123 ACCAGTGCCTCTCTCTCTGGCCCTCTTGTACAGGAGGAGAGCTGGCCCATCAGC 182  
Db 122 ACCAGTGCCTCTCTCTCTGGCCCTCTTGTACAGGAGGAGAGCTGGCCCATCAGC 181  
Qy 183 TCCCACTGCAGGCTTGACAAAGTCCAACTTCAGCAGCCCTATATCACCAACCGCACCTTC 242  
Db 182 TCCCACTGCAGGCTTGACAAAGTCCAACTTCAGCAGCCCTATATCACCAACCGCACCTTC 241  
Qy 243 ATCTGGCTAAGGAGG 258  
Db 242 ATCTGGCTAAGGAGG 257

RESULT 15  
US-09-870-574-1  
; Sequence 1, Application US/09870574  
; Patent No. 6551799  
; GENERAL INFORMATION:  
; APPLICANT: Gurney, Austin L.  
; APPLICANT: Aggarwal, Sudeepa  
; APPLICANT: Xie, Ming-Hong  
; APPLICANT: Maruoka, Ellen M.  
; APPLICANT: Foster, Jessica S.  
; APPLICANT: Goddard, Audrey  
; APPLICANT: Wood, William I.  
; TITLE OF INVENTION: INTERLEUKIN-22 POLYPEPTIDES, NUCLEIC ACIDS ENCODING  
; THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDERS  
; FILE REFERENCE: P2806-1(US)  
; CURRENT APPLICATION NUMBER: US/09/870,574  
; CURRENT FILING DATE: 2001-05-30  
; PRIOR APPLICATION NUMBER: US 60/169,495  
; PRIOR FILING DATE: 1999-12-07  
; PRIOR APPLICATION NUMBER: PCT/US00/14042  
; PRIOR FILING DATE: 2000-05-22  
; PRIOR APPLICATION NUMBER: PCT/US00/23328  
; PRIOR FILING DATE: 2000-08-24  
; NUMBER OF SEQ ID NOS: 7  
; SEQ ID NO 1  
; LENGTH: 1152

; TYPE: DNA  
; ORGANISM: Homo Sapien  
; US-09-870-574-1  
Query Match 5.1%; Score 244; DB 3; Length 1152;  
Best Local Similarity 100.0%; Pred. No. 1.1e-57;  
Matches 244; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 15 CTTCAGAACAGAGTTCTCTCTCCCTCCAGTCACCAAGTTCGAGTTAGAAATGTCTGCAATG 74  
Db 1 CTTCAGAACAGAGTTCTCTCTCCCTCCAGTCACCAAGTTCGAGTTAGAAATGTCTGCAATG 60  
Qy 75 GCGGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGAGCCCTGGCCACCAAGTGCCTC 134  
Db 61 GCGGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGAGCCCTGGCCACCAAGTGCCTC 120  
Qy 135 CTTCTCTTGGCCCTCTTGTGTACAGGAGGAGAGCTGGCCCATCAGCTCCCACTGCAGG 194  
Db 121 CTTCTCTTGGCCCTCTTGTGTACAGGAGGAGAGCTGGCCCATCAGCTCCCACTGCAGG 180  
Qy 195 CTTGACAAAGTCCAACTTCCAGCAGCCCTATATCACCAACCGCACCTTCATGCTGGCTAAG 254  
Db 181 CTTGACAAAGTCCAACTTCCAGCAGCCCTATATCACCAACCGCACCTTCATGCTGGCTAAG 240  
Qy 255 GAGG 258  
Db 241 GAGG 244

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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:16 ; Search time 2399.07 Seconds  
(without alignments)  
16534.837 Million cell updates/sec

Title: US-09-751-797-25

Perfect score: 4797

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Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 9793542 seqs, 4134689005 residues

Total number of hits satisfying chosen parameters: 19587084

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Published Applications NA Main:\*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	4797	100.0	4797	7	US-10-627-273-25
3	1349	28.1	1351	4	US-09-925-065A-79886
4	1095.6	22.8	1108	4	US-09-925-065A-687291
5	1073.6	22.4	1074	5	US-10-027-632-118181
6	1073.6	22.4	1074	6	US-10-027-632-118181
7	686	14.3	7445	3	US-09-751-797-8
8	686	14.3	7445	7	US-10-627-273-8
9	650	13.6	5935	3	US-09-751-797-29
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22	244	5.1	1152	3	US-09-870-574-1
23	244	5.1	1152	5	US-10-006-867-153

24	244	5.1	1152	5	US-10-066-500-125	Sequence 125, App
25	244	5.1	1152	5	US-10-063-547-153	Sequence 153, App
26	244	5.1	1152	5	US-10-063-551-153	Sequence 153, App
27	244	5.1	1152	5	US-10-063-616-153	Sequence 153, App
28	244	5.1	1152	5	US-10-063-569-153	Sequence 153, App
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32	244	5.1	1152	5	US-10-063-502-153	Sequence 153, App
33	244	5.1	1152	5	US-10-063-549-153	Sequence 243, App
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37	244	5.1	1152	5	US-10-066-494-125	Sequence 125, App
38	244	5.1	1152	5	US-10-230-163-243	Sequence 243, App
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41	244	5.1	1152	5	US-10-066-211-125	Sequence 125, App
42	244	5.1	1152	5	US-10-066-193-125	Sequence 125, App
43	244	5.1	1152	5	US-10-230-338-243	Sequence 243, App
44	244	5.1	1152	5	US-10-063-553-153	Sequence 153, App
45	244	5.1	1152	5	US-10-218-631-243	Sequence 243, App

ALIGNMENTS

RESULT 1

US-09-751-797-25

; Sequence 25, Application US/09751797

; Patent No. US20010024652A1

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Renauld, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; FILE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5543.2

; CURRENT APPLICATION NUMBER: US/09/751,797

; CURRENT FILING DATE: 2000-12-29

; PRIOR APPLICATION NUMBER: 09/419,568

; PRIOR FILING DATE: 1999-10-18

; PRIOR APPLICATION NUMBER: US09/178,973

; PRIOR FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 29

; SEQ ID NO 25

; LENGTH: 4797

; TYPE: DNA

; ORGANISM: Homo sapiens

; FEATURE:

US-09-751-797-25

Query Match 100.0%; Score 4797; DB 3; Length 4797;  
Best Local Similarity 100.0%; Pred. No. 0;  
Matches 4797; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	TGCACAGCAGAAATCTTCAGAACAGGTTCTCTCCCTCCACGTCACCACTGCTCGAGTTAG	60
Db	1	TGCACAGCAGAAATCTTCAGAACAGGTTCTCTCCCTCCACGTCACCACTGCTCGAGTTAG	60
Qy	61	AATTGCTGCAATGGCGCCCTGCAGAAATCTGTGAGCTCTTTCCTTATGGGACCTGG	120
Db	61	AATTGCTGCAATGGCGCCCTGCAGAAATCTGTGAGCTCTTTCCTTATGGGACCTGG	120
Qy	121	CCACGAGCTGCTCTCTCTGCGCTCTGTGTACAGGAGGAGGAGCTGCGCCCATCA	180
Db	121	CCACGAGCTGCTCTCTCTGCGCTCTGTGTACAGGAGGAGGAGCTGCGCCCATCA	180
Qy	181	GCTCCCACTGAGGCTTGACAACTTCCAGAGCCCTATATCACCACCGCACCT	240
Db	181	GCTCCCACTGAGGCTTGACAACTTCCAGAGCCCTATATCACCACCGCACCT	240
Qy	241	TCATGCTGGCTAAGGAGGTATACATCTCAATCTCTTCTTCTGTTGGATCTACTTGA	300



2461 QY CATTAAATGCAATGCTTTGAAACTTTGGAAGAAATAAACTCAGAA CAATGAGAAAAAGAGCTGG 2520  
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2521 QY ACTTGCATATAGGGCTAAATTTCTGGAGTAATAACACCTATTATTGAATTAATCATATATC 2580  
2521 Db ACTTGCATATAGGGCTAAATTTCTGGAGTAATAACACCTATTATTGAATTAATCATATATC 2580  
2581 QY TATCAGATATTGATTATAGTTTAAAGCAAGAGCAGACAACCCCGATCTCTTTTATACAG 2640  
2581 Db TATCAGATATTGATTATAGTTTAAAGCAAGAGCAGACAACCCCGATCTCTTTTATACAG 2640  
2641 QY GTTCAAAATAGAGTAAATAATATTAGTAAGAGATTATATTAGTTAAATGGAAGCTGGAATT 2700  
2641 Db GTTCAAAATAGAGTAAATAATATTAGTAAGAGATTATATTAGTTAAATGGAAGCTGGAATT 2700  
2701 QY GGTAAAGCTTTTTTCTCTCTCTCCCATCAAGACCTTCCATCTAGATTCTTCTCTTCA 2760  
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2761 QY CTCCTCAACAAATCCCTAGGGAGCATTTATCCATGGTGGCTGGTGTACATTTCTATAG 2820  
2761 Db CTCCTCAACAAATCCCTAGGGAGCATTTATCCATGGTGGCTGGTGTACATTTCTATAG 2820  
2821 QY TGAATGATACCATCATGTGGCTTATTTGGTGAAGAACAACAATGGAAGGCTTAGACTA 2880  
2821 Db TGAATGATACCATCATGTGGCTTATTTGGTGAAGAACAACAATGGAAGGCTTAGACTA 2880  
2881 QY ACAATAGTGAATCAACCAAAACCGGAGGAATGATTTAGGAGCAGTGAAGTGAGCTCTT 2940  
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2941 QY GCAAGCAGGTACCACTAAATCTCAGAAACATGAAGGCTCCAGTTGATGAAATTTTCAGT 3000  
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3421 Db GCAATGTTGTTTCAGAGAAAGATCAACAGAGGAGAAACTGTCTAGAGCTGTCTGAAATAGG 3480  
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US-10-627-273-25  
; Sequence 25, Application US/10627273  
; Publication No. US20040110189A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; APPLICANT: Louhed, Jamila  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: 2003-07-25  
; PRIOR FILING DATE: 2003-07-25  
; PRIOR APPLICATION NUMBER: US/10/627,273  
; PRIOR FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 25  
; LENGTH: 4797  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
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Query Match 100.0%; Score 4797; DB 7; Length 4797;  
Best Local Similarity 100.0%; Pred. No. 0;  
Matches 4797; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
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US-09-925-065A-79886  
; Sequence 79886, Application US/09925065A  
; Publication No. US20050228172A9  
; GENERAL INFORMATION:  
; APPLICANT: Wang, David G.  
; TITLE OF INVENTION: Identification and Mapping of Single  
; FILE OF INVENTION: Nucleotide Polymorphisms in the Human Genome  
; FILE REFERENCE: 108827.135  
; CURRENT APPLICATION NUMBER: US/09/925,065A  
; CURRENT FILING DATE: 2001-08-08  
; PRIOR APPLICATION NUMBER: US 60/243,096  
; PRIOR FILING DATE: 2000-10-24  
; PRIOR APPLICATION NUMBER: US 60/252,147  
; PRIOR FILING DATE: 2000-11-20  
; PRIOR APPLICATION NUMBER: US 60/250,092  
; PRIOR FILING DATE: 2000-11-30  
; PRIOR APPLICATION NUMBER: US 60/261,766  
; PRIOR FILING DATE: 2001-01-16  
; PRIOR APPLICATION NUMBER: US 60/289,846  
; PRIOR FILING DATE: 2001-05-09  
; NUMBER OF SEQ ID NOS: 957086  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 79886  
; LENGTH: 1351  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-09-925-065A-79886

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Qy 3607 AGATAGAAAGATATCTTTTGGCCCTATGTCAAAAGAGATATGAAAGGTGAAGGGC 3666
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Qy 3667 GGAAGAAAGCAGGAAAGGAACCATGTATTATATAGAGGCAATGGTGACAAGTTT 3726
Db 1201 GGAAGAAAGCAGGAAAGGAACCATGTATTATATAGAGGCAATGGTGACAAGTTT 1260
Qy 3727 TTCTTGAATAATGCAAAATATGATAGATTAGAGGAATTTTCAGTAGGGAATGCTTTTCACT 3786
Db 1261 TTCTTGAATAATGCAAAATATGATAGATTAGAGGAATTTTCAGTAGGGAATGCTTTTCACT 1320
Qy 3787 TGAATTTGGTTCCTCTTCGATTAAAGTTTG 3817
Db 1321 TGAATTTGGTTCCTCTTCGATTAAAGTTTG 1351
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RESULT 4

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US-09-925-065A-687291
; Sequence 687291, Application US/09925065A
; Publication No. US20050228172A9
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single
; FILE OF INVENTION: Nucleotide Polymorphisms in the Human Genome
; FILE REFERENCE: 108827.135
; CURRENT APPLICATION NUMBER: US/09/925, 065A
; CURRENT FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: US 60/243, 096
; PRIOR FILING DATE: 2000-10-24
; PRIOR APPLICATION NUMBER: US 60/252, 147
; PRIOR FILING DATE: 2000-11-20
; PRIOR APPLICATION NUMBER: US 60/250, 092
; PRIOR FILING DATE: 2000-11-30
; PRIOR APPLICATION NUMBER: US 60/261, 766
; PRIOR FILING DATE: 2001-01-16
; PRIOR APPLICATION NUMBER: US 60/289, 846
; PRIOR FILING DATE: 2001-05-09
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; NUMBER OF SEQ ID NOS: 957086
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 687291
; LENGTH: 1108
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-925-065A-687291

Query Match      22.8%; Score 1095.6; DB 4; Length 1108;
Best Local Similarity 99.8%; Pred. No. 3.1e-267;
Matches 1106; Conservative 1; Mismatches 0; Indels 1; Gaps 1;

Qy 1058 TCCTTCCACAGAGACCCCTTACCCCAACTCTCTCTCTTCCCTTACCCCTTACCTAGCTAGC 1117
Db 1 TCCTTCCACAGAGACCCCTTACCCCAACTCTCTCTCTTCCCTTACCCCTTACCTAGCTAGC 60

Qy 1118 AGGAAGAAGTGTCTTGGCAGCAGTGTATCAGAGAGTCATTTGGGATCATAGAGTATTTCG 1177
Db 61 AGGAAGAAGTGTCTTGGCAGCAGTGTATCAGAGAGTCATTTGGGATCATAGAGTATTTCG 120

Qy 1178 TTTTGTCTTTGACTGAGTCACTCTTCTGAGTTTATAGTGGTGAATGGGGTCTGGAACTTAAAG 1237
Db 121 TTTTGTCTTTGACTGAGTCACTCTTCTGAGTTTATAGTGGTGAATGGGGTCTGGAACTTAAAG 180

Qy 1238 TGTACAGAAAGCCGATTTGGTTTGTCTTCGAAAAAAGGCAACTCAGGTTCCGTAAGATGA 1297
Db 181 TGTACAGAAAGCCGATTTGGTTTGTCTTCGAAAAAAGGCAACTCAGGTTCCGTAAGATGA 240

Qy 1298 GAAAGTGTGGG-AAAACTCTAGCTGTGGAATGATCCATTTAGTCTTAAAGTTTGA 1356
Db 241 GAAAGTGTGGGAAAAAACAATCTAGTGTGGAATGATCCATTTAGTCTTAAAGTTTGA 300

Qy 1357 GGGAGGGGATGTCATGGAGAGAAATTTAGAAAGAAAGTGGGAAATGGGAAGGCTTAAAG 1416
Db 301 GGGAGGGGATGTCATGGAGAGAAATTTAGAAAGAAAGTGGGAAATGGGAAGGCTTAAAG 360

Qy 1417 TCGGTGGTGGGTGGCAGACTGTGTCCTCTGTTGATGTCTATGGGAAGCCACAAATCGGAG 1476
Db 361 TCGGTGGTGGGTGGCAGACTGTGTCCTCTGTTGATGTCTATGGGAAGCCACAAATCGGAG 420

Qy 1477 GCGTGTGAATTTGATCCGCTGNAACATTTGAAACTATGAAAAAGTTTTCAGTGGAGTGG 1536
Db 421 GCGTGTGAATTTGATCCGCTGNAACATTTGAAACTATGAAAAAGTTTTCAGTGGAGTGG 480

Qy 1537 GCCCAGTAAAAAGCCCTTAGGACTTCTAGAGAGGGCTTAATTTTCACTCAGATGTTTTA 1596
Db 481 GCCCAGTAAAAAGCCCTTAGGACTTCTAGAGAGGGCTTAATTTTCACTCAGATGTTTTA 540

Qy 1597 TGTACATTTTCTTGTCTTAAGCATGCAATTTTCTGGAGATACGATTTGAGGTTTTATTCCTT 1656
Db 541 TGTACATTTTCTTGTCTTAAGCATGCAATTTTCTGGAGATACGATTTGAGGTTTTATTCCTT 600

Qy 1657 ACAGAAATTTGCATAAACTACTCCGCTCTTTCACAAATGCCAAACCTCAGTAGGATTTCCC 1716
Db 601 ACAGAAATTTGCATAAACTACTCCGCTCTTTCACAAATGCCAAACCTCAGTAGGATTTCCC 660

Qy 1717 AAGATGAAGAGAGGTCCTTGTGAAGGAAGTCACTGGATTTCTGGGCTCCAAAGGAATTC 1776
Db 661 AAGATGAAGAGAGGTCCTTGTGAAGGAAGTCACTGGATTTCTGGGCTCCAAAGGAATTC 720

Qy 1777 AAGAGCTCAGGAAATCTAGGTCATCTGTGAAATCTAGGTCATTTGTGGGCAAAATTAATA 1836
Db 721 AAGAGCTCAGGAAATCTAGGTCATCTGTGAAATCTAGGTCATTTGTGGGCAAAATTAATA 780

Qy 1837 GAGCTTTAATTCAGGTGAATTTGATCTGTACCTCCATGGGTGGAGGTTTCATAAAGTTT 1896
Db 781 GAGCTTTAATTCAGGTGAATTTGATCTGTACCTCCATGGGTGGAGGTTTCATAAAGTTT 840

Qy 1897 CAGCAACAATTAAGATAGTTATGCTTTTATTTAGCATATTTGAAGTGTATGAC 1956
Db 841 CAGCAACAATTAAGATAGTTATGCTTTTATTTAGCATATTTGAAGTGTATGAC 900

Qy 1957 CTGCATATCCAGAGGAATGTGCAAAAGCTGAAGGACACAGTGAAGGAGGTAGGACTGATA 2016
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; PRIOR FILING DATE: 1999-11-23
; PRIOR APPLICATION NUMBER: US 60/156,358
; PRIOR FILING DATE: 1999-09-28
; PRIOR APPLICATION NUMBER: US 60/146,002
; PRIOR FILING DATE: 1999-08-09
; NUMBER OF SEQ ID NOS: 325720
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 118181
; LENGTH: 1074
; TYPE: DNA
; ORGANISM: Human
; US-10-027-632-118181

Query Match 22.4%; Score 1073.6; DB 6; Length 1074;
Best Local Similarity 99.9%; Pred. No. 1.2e-261;
Matches 1073; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2472 TGCCTTGAACCTTGGAGAATAAATCTCAGAACATGAGAAAGAGCTGGACATGATATA 2531
DB 1074 TGCTTTGAAACTTGGAGAATAAATCTCAGAACATGAGAAAGAGCTGGACATGATATA 1015
QY 2532 GGCTAAATTCGAGTAATAACACTTATTTTGAATTATCATAATATCTATCAGATATT 2591
DB 1014 GGGCTAAATTCGAGTAATAACACTTATTTTGAATTATCATAATATCTATCAGATATT 955
QY 2592 GATTATAGTTTAAAGCAAGCAGACAAACCCGATCTCTTTTATACAGGTTCAAAATAGA 2651
DB 954 GATTATAGTTTAAAGCAAGCAGACAAACCCGATCTCTTTTATACAGGTTCAAAATAGA 895
QY 2652 GTAAATATATAGTAAGATTTATATATAGTTTAAATGGAAGTCTGAATGCGTAACTTTT 2711
DB 894 GTAAATATATAGTAAGATTTATATATAGTTTAAATGGAAGTCTGAATGCGTAACTTTT 835
QY 2712 TTTTCTCTCTCCCATCAAGACCTTCCATTTAGTTTCTTCTTCTTCTTCTTCTTCTTCTT 2771
DB 834 TTTTCTCTCTCTCCCATCAAGACCTTCCATTTAGTTTCTTCTTCTTCTTCTTCTTCTTCTT 775
QY 2772 AATCCCTTAGGGAGCATTTATCCATGGTGGCTGGTGATACATTTCTATAGTAATGATACC 2831
DB 774 AATCCCTTAGGGAGCATTTATCCATGGTGGCTGGTGATACATTTCTATAGTAATGATACC 715
QY 2832 ATCATGTGGCTATTTGGTGAAGAAGAAACAATGGAAGGCTTAGACTAAACAATAGTGAC 2891
DB 714 ATCATGTGGCTATTTGGTGAAGAAGAAACAATGGAAGGCTTAGACTAAACAATAGTGAC 655
QY 2892 TCACCCCAAAACCGGAGGATGATTAGGACAGTGAAGTGACGCTCTTTCGACAGCGTAA 2951
DB 654 TCACCCCAAAACCGGAGGATGATTAGGACAGTGAAGTGACGCTCTTTCGACAGCGTAA 595
QY 2952 CAATAAATACTCAGAAACATGAAGGCTCCAGTTGATGGAATTTTCAAGTAAACAAGCTTAA 3011
DB 594 CAATAAATACTCAGAAACATGAAGGCTCCAGTTGATGGAATTTTCAAGTAAACAAGCTTAA 535
QY 3012 CCTTAATTCCTCTTTCCTCTGTACTTTTAAAAAAGCGTTTCTTCTGAGCAATCAT 3071
DB 534 CCTTAATTCCTCTTTCCTCTGTACTTTTAAAAAAGCGTTTCTTCTGAGCAATCAT 475
QY 3072 TTAATGAGTGATGCTGTTTCTTCTTGTATTAATGAAGGCTTTGTAGTTTAAATTTGTA 3131
DB 474 TTAATGAGTGATGCTGTTTCTTCTTGTATTAATGAAGGCTTTGTAGTTTAAATTTGTA 415
QY 3132 AGCCCAAGTCTCTGTTTATAGAACTATTATCTAGACATGAGGCGCTGAATGTTAGCATGC 3191
DB 414 AGCCCAAGTCTCTGTTTATAGAACTATTATCTAGACATGAGGCGCTGAATGTTAGCATGC 355
QY 3192 CACAGACAAGGCATGCTTTTACACATCTTGCTTAAAAAATTAAGTATTCATCTTGCTTGT 3251
DB 354 CACAGACAAGGCATGCTTTTACACATCTTGCTTAAAAAATTAAGTATTCATCTTGCTTGT 295
QY 3252 TGCTTTTAAAGAGTGAAGTGTGAGAGAGGAGATCTCATGTTGATCTGTGTGATTTTCA 3311
DB 294 TGCTTTTAAAGAGTGAAGTGTGAGAGAGGAGATCTCATGTTGATCTGTGTGATTTTCA 235

RESULT 7
US-09-751-797-8
; Sequence 8, Application US/09751797
; Patent No. US20010024652A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renauld, Jamila
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.2
; CURRENT APPLICATION NUMBER: US/09/751,797
; CURRENT FILING DATE: 2000-12-29
; PRIOR APPLICATION NUMBER: 09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 8
; LENGTH: 7445
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
; US-09-751-797-8

Query Match 14.3%; Score 686; DB 3; Length 7445;
Best Local Similarity 53.8%; Pred. No. 1.5e-162;
Matches 2644; Conservative 0; Mismatches 1875; Indels 393; Gaps 44;

QY 29 CTCCTTCCCGAGTCACAGTTGCTCGAGTTAGATTGTCTGCAATGSCCGCCCTGCGAGAA 88
DB 2034 CTCCTCTCTCACTTATCAACTGTTGACACTTGTGCGATCTCTGATGGCTGTCTGCGAGAA 2093
QY 89 ATCTGTGAGCTCTTTCCTTATGGGACCCCTGGGCCACCAGCTGCTCTCTCTTGGCCCT 148
DB 2094 ATCTATGAGTTTTCCTTATGGGACCTTGGCGCCAGCTGCTGCTCTCTCATTTGGCCT 2153
QY 149 CTTTGTACAGGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGCGAGGTTGCAAGTCCAA 208
DB 2154 GTGGGCCAGAGGAGCAAAATGCGCTGCCCGTCAACACCCGGTCAAGCTTGAAGTGTCCAA 2213
QY 209 CTTTCCAGAGCCCTATATACCAACCGCACCTTCATGCTGCTGCTGCTGCTGCTGCTGCTGCT 268
DB 2214 CTTTCCAGAGCCCTGATATCGTCAACCGCACCTTTATGCTGCGCAAGAGGTACAGCTGCA 2273
QY 269 AATCCTGCTCTTCTGTTGGATCTACTTGAATCCAAATAGTTCTTAAACTTTTCTTCA 328
DB 2274 TCTCTTCTCTCCATACCGCTTGGCAATTTTCTGAGGACCTTGCATACTTTTAGGG 2333
QY 329 GAGCATCTCTTAAGAGCTTTTAGGAAACCCACCTGTTTATCCCTGAGGGTAGATAAATTTCTG 388
DB 2334 CGCTTATCTCCGAGGCTCTCACTACCTAGTTT-----TCTGCTCTTTAGAG 2382
QY 389 TTTTTCAGAGACTCTTTGGGAATCTGGCTTTTTCCTTTTTCCTTGAACCTTCTTCCCTCAT 448
DB 2383 ACTCTTTAAGGACTGGGCTCTTTTCTATTTCTATTTCAAGGCTCTCAGGACCATTTCTTAT 2442

QY 449 TTTGGCCTTTATGATACATATGAAATTTTCCAAAGAGCGGCCATTTCAGTAATCCAT 508  
DB 2443 CTTGGCCCTTCAGGACATATACATACTGAATTTATCTACAGAGGCCATTT--AGNAGCCA 2500  
QY 509 CTGATGATTTTTTTTCCCTTTATGSCCTCTGTGCANTGTTCTAAACTCATGACACACATCTG 568  
DB 2501 CCCACGACTGCAATFACATTTTCCATTTCTGTGCTCTCTCTGMACTCATACTCTCTTTGGC 2560  
QY 569 AATTCTGCTTTAGTCTTTATGATGTTGCTCTGGGAGACGGGATGGGCACATGTTCTAT 628  
DB 2561 TACTC-----CTGAGACCCACTGCGGACATACATCTCTAC 2595  
QY 629 GTATAAATTTTTTTTCTATTGCTCAATGTCCAGACCTTAGTCTTTTCTTTCTCTCTCCAG 688  
DB 2596 TTACAGGCTTTTCTTCATCTCTCTGTGTCACCCAGGCACTTAGGGTTTTTC--TCTCTTTTCAG 2654  
QY 689 GCTAGCTTGGCTGATAACAACACACAGAGCTTGTCTCAATGGGGAGAAACTGTTCCACGGA 748  
DB 2655 GCCAGCCTTCAGATAAACAACACAGAGCTCCGGCTCATCGGGAGAAACTGTTCCGAGGA 2714  
QY 749 GTCAGTGTAGCTACAGTTGTGACGAACAGGGCCGTGTGCCGTCCATGGGTACTTTGGGGT 808  
DB 2715 GTCAGTGTAGTCTCTCACTGTGTATGAGCAGGGC-----TAGCTGCGGGAGCT 2761  
QY 809 GGTGGTGTATGATGCTTTAGGTCCTTATCCCTTATGACCCCTTCTGTTTCCCTTCCACCTGC 868  
DB 2762 GGTGGACCTCTGGGATAG-----TCTGACGTATGACCCCTGCTGCTCTTGTCTTACCTGC 2817  
QY 869 AGATGAGTGAAGCGCTGCTATCTGATGAAGCAGGTGCTGAACTTCACCCCTTTGAAGAAGTGC 928  
DB 2818 AGGCTAAAGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCACCCCTGGAAGACGTT 2877  
QY 929 TGTTCCTCTCAATCTGATAGTTTCCAGCCTTATATGCAAGGAGGTGGTGCCTTCTCGGCCA 988  
DB 2878 TGCTCCCCCAGTCAGACAGGTTCCAGCCCTACATGCAAGGAGTGGTACCTTTTCTGACCA 2937  
QY 989 GGCTCAGCAACAGGCTAAGCACATGTGTAGTTTTCAGCTCTCAGCCTATGCCCTACCTACCC 1048  
DB 2938 AACTCAGCAATCAGCTCAGCTCCTGTGTAGTGTGACTCTGGCTACCTATGCTCCTCTCT 2997  
QY 1049 CTCCTTCCTCTTCCACAGAGACCCCTTACCCTCAACTCTCTCTCTCTCTCTCTCTCTCTCT 1108  
DB 2998 CTTCTCTCTTCTTATTCAGTAAGAACCCGAGGTCTGCGCTCTCTCTCTCTCTCTCTCTCTCT 3057  
QY 1109 TAAGCTAGCAGGAAGAAGTGTCTTGGGACGAGTGTATCAGGAGTCA-----TTTGGG 1161  
DB 3058 GGAGGGCCTCAGCACCCACCATCATAGGCCACTTGAAATAGGTCACAAAGGCTTTGGC 3117  
QY 1162 ATCATAGAGTATTTGCTTTGCTTTGACTGAGTCACATCTTGTAGTTTATAGTGGTGAATG 1221  
DB 3118 TTCAATTGAGTAATCTTTGAGTTTGTATGAGTGAAGCTTTATTTGTTTTATCCATGGAA 3177  
QY 1222 GGGCTGGAACCTTAAGTGTACAGAGCGCCTATGGTTTGTCTTCGGAAAAAGCAATC 1281  
DB 3178 AGAATCAACTCAATCTGTAGGTGAGAAGATGTTGGGAAAGGAAAGGCGCTAGAT 3237  
QY 1282 A-----GGTTGCGTGAAGATGAGAAGGTGTTGG 1309  
DB 3238 AGAGAAACAGATCTGCTGAGTATAGTACTTATGCGGGAGCAGCGGGCGATATCCACTGA 3297  
QY 1310 GAAACATCTAGCTGTGGAATGATCCATTTGAGTCTTAAGTTGTTGAGGGAGGGGATGG 1369  
DB 3298 GTACAAGTACTTTGTGGGGAGAGAAATCCACTGAGTACAAGTACTTTGTTGGCATGGAGATC 3357  
QY 1370 CATGAGAGAAATAGAGAGAGAAATGGGAAATGGGAAGGCTTAAA----- 1415  
DB 3358 CACTGAGTACAAGTACTTTGTGGGGAGGGGATGGCACAGAGCAAAAGTTGAAGGAGG 3417  
QY 1416 -----GTCCGTTGGGTGCGGACACTGTTGCCCTGTTGATGTGATGGGA 1460  
DB 3418 AAGATGGAGAGGCCTCATGGTTGGGGTGTGAAGGTCACCTCTTTTCCATGTGATGGAG 3477

QY 1461 AGCCACAAATCCGAGGCGGTGTGAACCTTGTATGATCGCGCTGAACATTTGAAACTATGAAAAA 1520  
DB 3478 AGTTAAGAAAAACCACT--GTGTGAGTTTGTATGTTCTTTCAGACACCCCACTATGAACAT 3536  
QY 1521 AGTTTGTAGTGGAGTGGGCCCACTAAAGGCCCTTAGGACTTTACTTGAAGAGGGCTTAATTTT 1580  
DB 3537 ATCCACGAGGAGCGGCAGACTGTGGAGACCTGGGCATTTTAGGGA--AGCGCGGCTTTT 3594  
QY 1581 CACATGAGATGTTTTTATGTACATTTCTTTGTTCTTAAGCATGCAATTTTTCTGAGATACGAT 1640  
DB 3595 CACACGAGAACTTTATGCTCATCTCTTGTGCTACACTCCCACTTTGATGAGGTTTCAGC 3654  
QY 1641 TGAGGTTTTATTCCTTACAGATTTGCAATAAATCTACCTCCGCTCTTTTCCACAAATGCAAAAC 1700  
DB 3655 TCAGGTTTCTGTTCT-----ACCGTTCTTGTCTACTCTGGTGGAAAC 3693  
QY 1701 CTGAGTAGGATTTTCCAAAGATGAAGAGAGTCTCTTGTAAAGGGAAGTGAATGATTTCTG 1760  
DB 3694 TTCAGTAGGATTTCCCAAGACGAGGACAGCTCTCTCTGTAAAGGAGGGAACCTTGGATTTC 3753  
QY 1761 GCGTCCCAAGGGAATTCAGAGAGCTCAGGAAATCTAGGTCACTGTTGAAATCTAGTCAATG 1820  
DB 3754 GTGTCTTAGAGAAAGAAATAGCTCAGAGAAATCTAGGTCAAGTGAAATCTAGGTCAAGC 3813  
QY 1821 TGGGCAAAATTTACTTAAGAGCTTTAAATTCAGGTGAATTTGTAATCTGTACTCCATGGGTGTG 1880  
DB 3814 GGGCAAAATGACTGAACGCTCTATTTCCAGGTGAACGGTCAAGTCACTCAGATATACTG 3873  
QY 1881 GAGGTTCAATAAGTTTTCAGCACAAATTAAGATAGTTATGCTTGTATTTGTTTATAGCA 1940  
DB 3874 AGGTATTTGGGCTCCCAAGGATAGATTTCTGTTAGTGA--GTCTGCTTTTATTTTTCAGCA 3932  
QY 1941 TATTGAAGTGTAGTACCTGCATATCCAGAGGAAATGTGCAAAAGCTGAAGACACAGTGAA 2000  
DB 3933 CATCAGGGGTGACACACAGAAATCCAGAAAGATGTCAAGAGGCTGAAGAGACAGTGAA 3992  
QY 2001 AAAGGTAGGACTGATTAATCTGTAAGTCAATGCAATAGGAGAGACAAATTTGTTT 2060  
DB 3993 AAGGTACTATTGGCAAGCCACAATACTAAGCCATTCAGTAG--GAGACGTGGGGATTTTC 4050  
QY 2061 TTCTTTCTCTTTCTTTTCCCATCACTTTGTGATTTTTCCTTGAATTTCTCTTGAATTTCTTACCAC 2120  
DB 4051 TTTCTCTGCTTCCAGTCCCTTCTACTTTGTAACAATTTTATTTGACTTTGTCTACTACTCTG 4110  
QY 2121 GGGCATTA---CTTTGCTGTGTGTATGTAGATATATCTATATATCTAGATGTCAGTT 2176  
DB 4111 GTCCATTTCTGCTTTAGTGCACCTGTATCTAGCTGGGTCTATAGATCTTTCAATCTGTG 4170  
QY 2177 TCCAAATCTTGCATAATTTGTAGAATTTAGAACTGGTTGGGATCTTAGCTTGTCTAGTCAAC 2236  
DB 4171 TCTAAATTT--GTAAGTCAAAATTTCTGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTC 4227  
QY 2237 ATAACTCTCAGATTTCTGGGGATGTCAGTGGCAGAGATAGGGCTAGATAGCAGGTCTCTCTG 2296  
DB 4228 ATGAGCACTTGTCTGGAGGATGGCTTGTGACAGAGTCAATGCTAGAAAGACAGCATCCCTG 4287  
QY 2297 AATCCCAAGCCACACTTTTCCCGTGTGATACAGATTTAGTTTGTGTTGTTGTTGTTGTTGTTGTT 2356  
DB 4288 ATTTCCAGCTGTCAC--TTGCTTAGTGGCCATGTTAATTTACTTTTGGCTTGAATAGTAT 4346  
QY 2357 TAGGAAATTTTCAGATTTCTTATTCACCTCATGTAATCTGAAGAAGTACTTTGTTTAAAAACA 2416  
DB 4347 TTGGGAAA--GCCAGTTCCCAAGCCCTACATATCTGAAGAACCATGCAATTTGAAGAACTA 4404  
QY 2417 GAAAAATGCTTATGGGCAAAATTTTATTTGAAGTCAATTTTGAAGTCAATTAATGCAATTTGCTT 2476  
DB 4405 GAAA---GCTGGGCAAAACTTACTAGAGATGATTTTGTGAGTCAATTTAAACGGATGCTC 4460  
QY 2477 TGAACTTTGGAAGATAAACTCAGAACAAATGAGAAAAAGAGCTGGACTTGCATATAGGGCT 2536  
DB 4461 TGAATTTGGGCAAAATCAACCCAGAAATAAACAAAAGAGCTGGATTTTGCAAATAGGACA 4520  
QY 2537 AATTTCTGGA-----GTAATAAACACTTTATTTTGAATTTATCATATAATCTATCATAGATA 2589







Db 3537 ATCCAGGAGCGGCGAGACTGTGGAGACCTGGCAATTTAGGGA--AGGCGGCGCTTTT 3594  
Qy 1581 CACATGAGATGTTTATGATACATTTCTTGTCTAAGCATGCAATTTCTGGAGATACGAT 1640  
Db 3595 CACAGAGAACTTTATGCTCATCTCTGTGCTACACTCCACCTTTGATGAGGTTGAGC 3654  
Qy 1641 TGAGGTTTATTCCTTACAGAAATTTGCAATAAATPACTCCGCTCTTTCCAAATGCAAAC 1700  
Db 3655 TCAGGTTTCGTTTCT-----ACCGTCTCTGCTACTGCTGGTGAAC 3693  
Qy 1701 CTGAGTAGGATTTCCCAAGATGAGAGAGTCTCTTGTGAAGGAGTGAATCTG 1760  
Db 3694 TTCAGTAGGATTTCCCAAGACGAGGACGCTCTTCTGTGAAGGAGGACCTGGATTTCA 3753  
Qy 1761 GCGTCCAAGGGAATTCAGAGCTCAGGAATCTAGGTCACTGTTGTAATCTAGGTCATTG 1820  
Db 3754 GTGTCTTAGAGAGCAATAGCTCAGAGAACTAGGTCAAGTCAAGTCTAGGTCACAGC 3813  
Qy 1821 TGGCAAAATTAACAGAGCTTTAAATCCAGGTGAATGTTACTGTACTCTCATGGGTG 1880  
Db 3814 GGGCAAAATGACTGAACGCTCTATTCAGGTGAACGGTCACTGCTCAGATATACTG 3873  
Qy 1881 GAGGTTCAATAAGTTTCAGCACACATTAAGATAGTTATGCTTGTATTCTTTTATAGCA 1940  
Db 3874 AGGTATTGGGCTCCACCGGATAGATTTCTGTTAGTGA-GTCTGCTTTTATTTTGCAGCA 3932  
Qy 1941 TATTGAAGGTGATGACCTGCATATCCAGAGAAATGTGCAAAAGCTGAAGCACACAGTGAA 2000  
Db 3933 CATCAGCGGTGACACAGCAACATCCAGAGAAATGTGCAAGGCTGAAGGAGACAGTGAA 3992  
Qy 2001 AAAGGTAGGACTGATACTGTCAATGTCAATGCTAATGCAATAGGAGAGACAAATGTTGTTT 2060  
Db 3993 AAAGGTACTATTGGCAAGCACAAATACATAAGCCATTCAAGTAG--GAGACGTGGGGATTTC 4050  
Qy 2061 TTCTTTCTTTCTTTCTCCATCACTTTGTGATTTTCACTGATTTCTCTACCACAG 2120  
Db 4051 TTTCTCTGCTTCCAGTCCCTTCTACTTTGTAACTTTTATTTGACTTGTCTACTACTG 4110  
Qy 2121 GGGCATTA---CTTTGGTGTGTGTATGTAGATATCTATATATCTAGATGTCAGTT 2176  
Db 4111 GTCAATTTCTCGTTAGCTGACCTGTATCTAGCTGGGTCTATAGATCTTTCAATCTGTG 4170  
Qy 2177 TCCAAATCTTGCAAAATGTAGAAATTCAGAACTGGTTGGGACTTAGCTTTGTCTAGTCAC 2236  
Db 4171 TCTAAATTT--GTAAGTCAAAATCTCTGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTC 4227  
Qy 2237 ATAACTCAGATTTCTGGGATGGTCAGTGCAGAGATAGGGCTAGAAATGCAAGTCTCCTG 2296  
Db 4228 ATGAGCACTTGTCTGGGAGATGGCTTGTGACAGAGTCAATGTCTAGAAAGACAGCATCCCTG 4287  
Qy 2297 AATCCCAAGCCAGCACATTTCCGGTGGTGATACAGATTAGTTTGGTACCATTAATTTCT 2356  
Db 4288 ATTCCCAAGCTCTGCAC--TTGCCCTAGTGGCCATGTGTAATTTACTTTGGCTTGTATAGTAT 4346  
Qy 2357 TAGGGAATTTTCAGATTTCTATTGACTCATGTAATCTGAAGAAGTACTTGTTTTAAACA 2416  
Db 4347 TTGGGAAA--GCCAGTTCCACGGAACCTACATAATCTGAAGAACCAATGCAATTTGAAACTA 4404  
Qy 2417 GAAAAATGCCATTTGGGCAAAATTTATTTGAAGTCAATTTTGAAGTCAATTAATGCAATGCTT 2476  
Db 4405 GAAA-----GCTGGGCAAAACTTACTAGAGATGATTTTGGAGCTCATTAACCGGATGCTC 4460  
Qy 2477 TGAACCTTGAAGAAATAAACTCAGAACAAATGAGAAAGAGCTGGACTTGCATATAGGGCT 2536  
Db 4461 TGAATGTGGCAAAATCAACCCAGAAATPACACAAAAGAGCTGGATTTGCAAAATAGGACA 4520  
Qy 2537 AATTTCTGGA-----GTAATAAACACTTATTTTGAAATTTATCATATATCTATCAGATA 2589  
Db 4521 AGTATTTAGAAATCACTGTTATTAATAGCTATCATCTTAAATTAATAATATAGGCTTATATA 4580  
Qy 2590 TTGATTTATAGTTTAAAGCAGAGCAGACAAACC--CCGATCTCTTTTATACAGGTTCAAT 2648

Db 4581 TATATTTAAGATTAACACAAAGAGTGGATAGCCCTCCCAATTTACTTGGCCTGGTTTCAAA 4640  
Qy 2649 AGAGTAAAAATATTAGTAAGAGATTTTATTAGTTAAATGGAAGTCTCAATTTGTAAGCT 2708  
Db 4641 AGAGTAAAAATATACAGTCATGGATTAATATTAGTGTATGATGATGAGATGGAACCC 4700  
Qy 2709 TTTTTCCTTCTCTCTCCCATCAAGACCTTCCATTTAGTTCTTCTTCTTCACTTCCCTCA 2768  
Db 4701 TTTCTTACTTTTACCTTCA-----TTTCTTAGTTTCTTTTCTTCTTCAACCCCTGA 4752  
Qy 2769 ACAAATCCCTTAGGAGCATTTATCCATGTGGCTGGTGTACATTTCTATAGTGAATGAT 2828  
Db 4753 TCAAGCCACTAGTAAGACCTATCTGCTGTGAGCTATTATATGACTTTTACAGCAACAC 4812  
Qy 2829 ACCATCATGTGGGCTATTTTGGTGAAGAAACA--ACAATGGAAGGCTTAGACTAACATA 2886  
Db 4813 ATTGCTGTGTGGCTCTTTTGGGAAGGAACAGATAGCAGGAGCTCAGGCTAGCAAGT 4872  
Qy 2887 GTGACTCACCCCAAAACCGGAGGAATGATTAGGAGAGTGAAGAGTACGCTCTTT--GCAAG 2945  
Db 4873 CTGACTTGGCCTTAAAGCCAGAGCATGGTTTATAGCAGAGAAAGTGAGGCTCTTTCGCAAG 4932  
Qy 2946 CAGGTACAACTAAATACTCAGAAACATGAAGGCTCCAGTTGTAGTGAATTTTTCAGTAACA 3005  
Db 4933 TGGGTGTGTTAAGTAATCAGAAACAGGAAGGCTCCGGTTGATGGAATTTATCAGTAAGAT 4992  
Qy 3006 GCTTAACCTTAATTTCCCTTTTCCCTCTTGAACCTTTTAAAAAAGCGTTTCTTCTCTGAG 3065  
Db 4993 ATCTACCTTATCTCTTCTATCGAAC-----TAAATCGTCTCTTTTCTTGTG 5042  
Qy 3066 CATCATTTAATAGTGTGACTGTTTCTTCTTTGATAAATGAAAGGCTTTGTAGTTTAAA 3125  
Db 5043 TGTAGGCTGTATAAACACACTTGT--TTCTTTTGTAGTGTTCATGGCTTGTAGATTTTAA 5100  
Qy 3126 TTGTGAAGCCAGTTCTCTTGTATTAGAACTATTATCTAGACATGGAGGCTGAATCTTA 3185  
Db 5101 GTGCTCTGCGAGTCTTGTGTAGAG--GGTTGTGTACCTTGACACCTGGGCTGGATGTTA 5158  
Qy 3186 GCATGCCACAGACAGGATGCTTTTACATCTTGTCTTAAAAAATTAAGTATTCATCTT 3245  
Db 5159 GCATGCCAAGGCACACACTTCTGNAATGCCGTGTGTAAGAGTTATTAATCATTTACT--- 5215  
Qy 3246 GCTTGTGTCTTTTGAAGAGTGAAGTGTGAGAGAGGAATCTCATGGTGA----- 3296  
Db 5216 ----TTGTCTTTGGAAGGTGAAGCGTGTGTGAGAAAGAACTCACAGGAGATGTGTCT 5270  
Qy 3297 -----TCTGTGTGATTTTCAAGACTTTTAATCCATTTTGAAGATCAATTT 3342  
Db 5271 CTGTAGAAACTTTTTCCTTAAATGCTTAAATCCACTTTTCAAGTCAAC---TT 5327  
Qy 3343 TCATATTTGCAATGGGTTGCCATGTGGAAGAGTATTATGCTTTTGTGGTGGTGTCA 3402  
Db 5328 TGACTTTTATACCATGCTGTCAATGAAAGAGTGTTTAGGCCGCTCTCATGGCTCTGGG 5387  
Qy 3403 GAAAGCA--CAGGAGGAGAGCAATTTGTTTCAGAGAAAGATCAACAGGAGAGAGAACTGT 3461  
Db 5388 AAAAGCAACCAATAGGCGAAGGAATGTTATGCTGAGAAATCTGACCGGACAGGAACTGGT 5447  
Qy 3462 CAGAGCTGTGAAATAGGCTGTTTGGGAGGCAATTAATCCCTCTCTGTTGGGGTAAA 3521  
Db 5448 CAGAGCTCCCCCAAGACCA-----CCACAGGTTTAAAGTAGG 5485  
Qy 3522 AGCAGAAACGAGGTTGGTAGTAAAT--GCATGACAGACAGTAGGGGACGATTAACCTTAA 3580  
Db 5486 AACAGTCCAGGTTGGGCTCATGTAAATAGATGAAACAGCGAGGAGGAATAGACTACAA 5545  
Qy 3581 AATTCCTTTATGCTTTGGAGTCTTTGAGATGAAAAAGAAATATCTTTTGGCCTTATGCTCA 3640  
Db 5546 AGTTTCATAGGGTC--CGAGTCTTAAAGATACAAATAGCTGC--TTGGGCTTCATAACA 5602  
Qy 3641 AAAGAAGTATGGAAGG-----TGAAAGGGCGGAAGAAAGCAGAGAAAGGAG 3688  
Db 5603 AAGGAGTCTGGGAAGGCAGCAAGTAGAGAGGGAATTGAAGGGGAAAAAACAAGATCTAG 5662



Qy 3689 AACCATGTATTATATAGAGGCAATGCTGACAAAGTTTCTTCTGAAATAATGCAAAATATG 3748  
Db 5663 AGACCTTGAACAGCTCAAAATCTCTACCAAGCAATTTTCTTGGAAACAATCTAGAGGT 5722  
Qy 3749 ATAGATTAGAGGAATTTCACTAGGAAATGCTTTTCACTTGAATTTGGGTTTCCCTCT--T 3805  
Db 5723 AGTGAATTAGGTGATTGACGGGGACCTTCTTTGGCCATTTGAAATCTGGGTTTGTCTCT 5782  
Qy 3806 CGATTAAAGTTGGATCCTCATCTGCAATTTGACT----TGGAGAGAGAAAGATGAATGT 3861  
Db 5783 CCATTGAGTTGAAGGCTCACCTTTTACCCCTCGAATGAGGAGGAGAAAGAGGGGTGT 5842  
Qy 3862 TAGGACCTATATCTGGTTTCTATTAACTAAAGCAAGTGGAAAGACTTATTATTGGTATTT 3921  
Db 5843 TATGACTCTCCACCTGGAGTTTACTAGTTTACGCATTTGGAACACAGACACTCGGACCTCCT 5902  
Qy 3922 TTCCACAAAAGTGAAGACTTTCTTTTACTGTTTGTCAAAAAGGTGGAATAGAAAAG 3981  
Db 5903 CTTGACAAAAAATGGAACCTGTGTGTCTTGTCTTTTGTCTTTTGTAAAGAAAGCAC 5962  
Qy 3982 CCTTAATGTATGTGTAATACATCGTTCAAAGTCATTTGAGTAGAGATGTTTTAAATCAG 4041  
Db 5963 AGCAAAAGCCGACCAATGAGTTGAATGTGGGTCTTTTGAAGTCAAGGCTTTTGAGTTGAG 6022  
Qy 4042 GAGTGTCCAATCATTTGGCTTCCCTGGACCACTTGAAGAATTTGTCTTGTATCACACAT 4101  
Db 6023 CACTCATCAATAGTT-----GATCATGTGTCAGGTGGAGGGC 6058  
Qy 4102 AAAATACAAGAAACAATAGCTGATGAGCTTAAAAAGTCCATGCAATAATCTCATACTGTTT 4161  
Db 6059 TACCTGTGACGCCGAGCCCTGCTGCTTGGCACATTAACATCTCCAGGTCTCAGTATCACT 6118  
Qy 4162 TAAGAAAGTTTATGAATTTCTGTAGGTGCAATTTCAAAGCTGTCTCTGGGCCATGTGGGC 4221  
Db 6119 TCCTGCTACTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTCCT----- 6163  
Qy 4222 CTGTGGGCTGCAGGTTGGACAAGCTCCTTTAAGTAATCTGTCTATAGATAGTTTGGAGC 4281  
Db 6164 -----AACCCCACTAAATTTAATTTGACAAAGACTGTGTAATTTG 6205  
Qy 4282 TGCAAAACAGGCCCAAGGCAATAGGTGGGCACCTCGGGATCCCCAGATGCCAGGCTCACT 4341  
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Qy 4342 TCAGTCTCCTTGTCTGTGTTAAGAAAGGGTGGTCAACTCTCTGCCAGCTTTTAAACAGC 4401  
Db 6228 ----TCTATGTGTGCAATTTGTCAAGGTTTCAATAAGATAGATTAAATAGGCCCATCAACAGC 6283  
Qy 4402 TTCATTAGTGTGAGGTGCACCTGAAATTTGATGCTGTGTGGTGGCCT-CTCAGTCCAGAGA 4460  
Db 6284 TTTATGGGTGGAATGCAAGTAATATAGGTAGATGCTGTGGTGTCTTATAGGTCAAGAA 6343  
Qy 4461 GCCGTCAATTTAAGCTCTTTGGCAAAATCATACAATCTATAAGGGATA-----T 4508  
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Qy 4509 TACTATGAATGTTTACAAATGCTTAAATCTCGGTTCTGTCTCCATCAACCTTAATCTTTG 4568  
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Qy 4569 CAATTTCT--AAATTTGTTCACTTTAGAAACATGCGCATTAATGCTCAAAATCTTTTGA 4625  
Db 6464 CCAGCTCTTTCTAATCTGTATCACTTTTAGAAAATGCTACTCTGTGCTCAAAATGGTTTGTGA 6523  
Qy 4626 TTCTTATTTTTCACAGCTTTGGAGAGTGGAGAGATCAAGCAATTTGGAGAACTTGGATTG 4685  
Db 6524 TTCATTATTTTCATAGCTTTGGAGAGTGGAGAGATCAAGGCGATTGGGGAAGCTGGACCTG 6583  
Qy 4686 CTGTTTATGCTCTGAGAAATGCTTGCATTTTGAACAGAGCAAGCTGAAAATGAAATAC 4745  
Db 6584 CTGTTTATGCTCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGGAAGAAC 6643

Qy 4746 TAACCCCTTTTCCCTGCTAGAAAATAACAATAGATGCCCCAAAGCGATTTTT 4797  
Db 6644 TGCTCCTTCTCGCTTCTTAAAGAAACAATAAGATCCTCGAATGGACTTTTT 6695  
RESULT 9  
US-09-751-797-29  
; Sequence 29, Application US/09751797  
; Patent No. US20010024652A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIPS) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/751,797  
; CURRENT FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 29  
; LENGTH: 5935  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-09-751-797-29  
Query Match 13.6%; Score 650; DB 3; Length 5935;  
Best Local Similarity 56.5%; Pred. No. 1.9e-153;  
Matches 1863; Conservative 0; Mismatches 1285; Indels 152; Gaps 29;  
Qy 29 CTCCTCCAGTCACAGTTGCTCGAGTTAGATTCTGCAATGCGCGCCCTGCGAGAA 88  
Db 356 CTCCTCCCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGGTCTCTCGCAGAA 415  
Qy 89 ATCTGTGAGCTTTTCTTATGGGAGCCTGGCCACAGCTGCTCTCTCTTTGGCCCT 148  
Db 416 ATCTATAGATTTTCCCTTATGGGACTTTTGGCCGACAGTGGCTTCTCATTTGCCCT 475  
Qy 149 CTTGGTACAGGAGGAGCAGCTGCCCATCAGCTCCCACTGCGAGGTTGCAAGTCCAA 208  
Db 476 GTGGGCCCGAGGCAAAATGCGTGCCTCAACACCCGGTGCAAGCTTGAGGTGTCCAA 535  
Qy 209 CTTCCAGAGCCCTATATACCAACCGACCTTCTGCTGCTAAGGAGGTATATCTC 268  
Db 536 CTTCCAGAGCCGTACATCGTCAACCCGACCTTTATGCTGGCCAAAGGAGGTACAGTCA 595  
Qy 269 AATCCTGCTCTTCTGTTGGATCTACTTGGAAATCCAAATAGTTCTTAAACTTTTCTCA 328  
Db 596 TCTCTTTCTTCCATACCGCCTTGCCTTCTCTGAGGACTTGCAAACTCTTTAGGGGC 655  
Qy 329 GAGCATCTTAAGAGCTTTTAGGAAACCACTGTTTATCCCTGAGGGTAGATAAATTTCTG 388  
Db 656 GCTTTATCTCCGAGGCTCTACTACTATGTTTCTGTCT-----CTTTAGAG 703  
Qy 389 TTTTTCAGAGACTCTTTTGGGAATCTGGCTTTTTTTTTTCTTGAACCTTCTCTCTCCAT 448  
Db 704 ACTCTTTAAGGACTGGATCTTTTCTATTTCTATTCTTCAAGGTCTCAGGACCAATTTCTTAT 763  
Qy 449 TTTGGCCTTTATGATACATATGATGAATTTTCCAAAGAGCGGCATTCAGTAAATCCAT 508  
Db 764 CTTGGCCTTTAGGACACATATACATGAATTTTATCTACAGAGCGGGTTT--AGAAAGCCA 821  
Qy 509 CTGATGATTTTTTTTTTCTTTATGCTCTGTGCAATTTGTTTAAACTATGACACATCTG 568  
Db 822 CCCAGACTGCAATACTTTTCCATCTGTGTGTCTCTTCTGAACTCATCTCTCTTGGC 881  
Qy 569 AATTCTGCTTTTAGTCTTTTATGATGTTGCTCTGGGAGACGGGATGGGGCAGATCTCTAT 628  
Db 882 TACTC-----CTGAGACCCCATGCGGACATACATCTCTAC 916





1222 GGGTCTGGAACTTAAGTGTACAGAGCGCATTTGGTTTGTCTTCGGAAAGCAACTC 1281  
1499 AGAAATCAACTCAAACTCTGAGATGAGAAAGATGTTGGGAACGAAAGGCGCTAGAT 1558  
1282 AGGTTGGCTAA--GATGAGAAAGGTTGGGAAACATCTAGCTGTGGAAATGGATCCA 1338  
1559 AGAGAAACAGATCTGCTGAGTACAGTACTTATGCGGGGGGGGCGAGGGCGGATATCCA 1618  
1339 TTGAGTCTAAGTCTGTCGAGGGGAGCGATGCGATGGAGAGAAATAGAGAGAAAGTGGG 1398  
1619 CTGAGTCTCAAGTACTTGTGGGAGAGAAATCCACTGAGTACAGTACTTGTGGGGGAAG 1678  
1399 AAATGGGAAGGCTTAAAGTGGTGGTGGTGGGAGACTGTTGCC-----TGTGA 1450  
1679 AATGGCACAGAGCAAAAGTTGAAGGGGAAAGAGGAAGATGAGAGGCGCTCAATGTTGGGG 1738  
1451 TGTATCGGGAAGCACAATAATCGAGGCGGTGTGAATGATGCCGCTGAAACATTTGAAC 1510  
1739 TGTGAAGGTCACCTCTCTTTTCCATGTGTGAGAGTTAAGAAATAACAGTGTGTGAGTT 1798  
1511 TATGAAAAAAGTTTGAAGTGGGCGGCCAGTAAAGGCCCTAGGACTTACTGAAGAGG 1570  
1799 TGATGCTTCAGACACCCCAACTATGCGAGCTGTGGAGACCTGGCAATTTAGGGA--AGG 1857  
1571 GCTTAAATTTTACATGAGATGTTTTATGTACATTTCTTGTCTAAGCATGCAATTTTCTG 1630  
1858 CGCGGCTTTTACACAGAGAACTTTATGCTCATCTCTTGTGTACACTCCACACCTTTGAT 1917  
1631 GAGATAGATTTAGGTTTTATCTCTTACAGAAATTTGATCAAACTACTCCGCTCTTCCAC 1690  
1918 GAGGTTAAGCTCAGTTTTGTTTTCT-----ACCGTTCTTGTCTAC 1956  
1691 AAATGCAAACTCAGTAGGATTTCCCAAGATGAAGAGGCTCTCTGTGAAGGAAGTGA 1750  
1957 TGGTGAACCTTCAGTAGGATTTCCCAAGACGAGGACGCTCTCTGTGAAGGAGGAGAC 2016  
1751 CTGGATTTCTGCGCTCAGAGGAATTTCAAGAGCTCAGGAAATCTAGGTCACCTGTTGAATC 1810  
2017 CTGGATTTCAAGTCTCAGAGAGCAAAATAGCTCAGAGAACTAGGTCACGTCMAATCT 2076  
1811 TAGTCAATTTGGGCAAAATTAAGAGCTTTAAATCCAGGTGAATTTGACTGTACCTC 1870  
2077 AGGTCACAGCGGGCAAAATGACTGAACGCGCTCTAATCCAGGTGAACGCTCACTGCGCTC 2136  
1871 CATGGGTGTGAGGTTTCATAAAGTTTCAGACACACATTAAGATAGTTATGCTGTTATTG 1930  
2137 AGATATCTCAGGTAATTTGGGCTCCACCGGATAGATTTCTGTTAGTGA-GTCTGCTTTTA 2195  
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2196 TTTTGCAGCACATCAGTGGTGACGACCAGAACATCCAGAGAAATGTCAGAAAGCTGAAG 2255  
1991 ACAGAGTAAAGGTAGGACTGATATACTGCTCAATGCTAAGTCAATGCAATAGAGAGACA 2050  
2256 AGACAGTGAAGAAAGGTACTATTGGCAAGCCACATACTAAGCCATTGAGTGGAGACGTG 2315  
2051 AATGTTGTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2110  
2316 GGGATTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2373  
2111 CTACCACAGGGGGAAT-----ACTTTGGTGTCTGTGTATGTAGATATATCTATATATCTA 2166  
2374 CTACTGTCTGTCCTCACTTACTCCTTACTGCTAGCTGCACTGATCTAGCTGGTCTATAGATCTT 2433  
2167 GATGTCAGTTTCCAAATCTTGGCAATTTGTGAATTTCTAGAACTGGTTGGGATCTTAGCTT 2226  
2434 TCAATCTGTGTCTAAATTT--GTAAGTCAAAATTTCTGGAGCTAGCAGAAAGCTTAGCTC 2490  
2227 GTCTAGTCAATACCTCAGATTTCTGGGATGTCAGTGGCAGAGATAGGCTTAGATGC 2286  
2491 AGCCAGTCTCATGACCTATGCTCGGAGGATGGCTTGTGACAGAGTCAATGCTAGAGAGAC 2550

## RESULT 11

US-10-027-632-208140  
; Sequence 208140, Application US/10027632  
; Publication No. US20020198371A1

2287 AGGTCTCTGAATCCCAAGCCAGCACCTTTTCCGGTGGTGATACAGATTAGTTTGGTAC 2346  
2551 AGCATCCCTGATTTCCAGCTCTGCAC--TTGCCTAGTGGCCACGTTGTAATTTACTTTAGCCT 2609  
2347 CATTAATTTCTAGGGAATTTTCAAGATTCCTATTGACTCATGTAATCTGAAGAGTACTTG 2406  
2610 GATTAAGTATTTGGGAAA--GCCAATTTCCACCGACCTACATAATCCGAAGAGCATGCA 2667  
2407 TTTAAAAACAGAAAAATGCTCATGCGCAAAATTTATTGGAAGTCATTTTTCGAAGTCATTAA 2466  
2668 TTGAATACTAGAAA-----GCTGGGCACAACTTACTAGAGATGATTTTGGAGCTCATTA 2723  
2467 TGCATTGCTTTGAACTTGGAGAAATAAATCTAGAAATAATGAGAAAGAGCTGGACTGC 2526  
2724 ACTGATGCTCTGAAATGTGATCAATCAACCCAGAAATAACAACAAAGAGAGCTGGATTTGC 2783  
2527 ATATAGGCTTAATTTCTGGAGTAAATAACACTTAT-----TTTGAATATCATATA 2578  
2784 AAATAGGACAAAGTATTTAGAAATCACTGGTATTAACAGCTGTCATCTTAATTTAAATATAG 2843  
2579 TCT---ATCAGATATTTAGTATATATAGTTTAAAGCAAGAGCAGACAAC--CCGATCTCTTT 2634  
2844 TGTCTATTTAGCTGCCTATTTAAGATTTAAACAAGAGTGGATACTTCCCAATTTACTG 2903  
2635 ATACAGGTTCAAAATAGAGTAAAAATATTAGTAAGAGATTTATTATAGTTAAATGGAAATC 2694  
2904 GGCCTGGTTTCAATAGAGTAAAAATATCAGTCATAGATTAATTTATAGTGTGATGAGAAATG 2963  
2695 TGAATTTGGTAAAGCTTTTTTTTCTTCTCTCTCTCCATCAAGACCTTCCATTTCTAGTTTCTT 2754  
2964 TGAGTTGGAAAAAC---CTTTCTCTTACTTTTTTACTCTTCTTCTTATTTATTTTCTTTT 3020  
2755 CTTTCTACTCCCTCAACAAATCCCTAGGAGCATTTATCCATGTGGGCTGGTGACATTT 2814  
3021 TCTTCAACCTCTGATCAAGCCACTAGTAGCACCTATCTCTCGAGCTATTATATGACT 3080  
2815 CTATAGTGAATGATACCATCATGTCCTCTATTGTTGGTGAAGAAACA--ACAATGGAAAGC 2872  
3081 TTACAGCAAAACAATTTGCTGTGTGGCTCTTTTGGGGAAGGGAACAGGATAGCAGGAGGC 3140  
2873 TTAGACTAACAA--TAGTGACTCACCCCAAAACCGAGGAATGATTAGGAGCAGTGAAGT 2931  
3141 TCAGGCTAGCAAGTCTGGACTCAACCTAAAGCCAGAGGCACTGGTTGATAGCAGAAAGT 3200  
2932 GAGCTCTT--GCAAGCAGGTACAACTAAATPACTCAGAAACATGAAGGCTCCAGTTGATGG 2990  
3201 GAGCTCTTCAAGAGTGGGTGCTTAAGTAATCAGAAACAGAGGCTCTGGTTGATGG 3260  
2991 AATTTTCAGTAAACAAGCTTAAACCTTAATTTCCCTTTTCTTCTTCTTCTTCTTCTTCTTCT 3050  
3261 AATTAATCAGTAAGATATCTACCTTATCTCC-----TTCTTCTATAGAAGCTAAACCG 3313  
3051 GCGTTTCTTCTGAGCATCATTTAATGAGTGTGACTGTTTCTTCTTCTTCTTCTTCTTCTTCTTCT 3110  
3314 TCTCTCTTCTTCTGTTGTGTAGGCTGATAAACACACGCTTGT--TTCTTCTTGTAGTGTCTAGG 3371  
3111 CTTTGTAGTTTTAAATTTGTGAAGCCAGTTCTCTGTTTATAGAACTATTATCTAGACATG 3170  
3372 CTTTGCAGATTTTCAAGTGTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 3429  
3171 GAGGGCTGAATGTTAGCATGCCACAGAGGCGATGCTTTTACACATCTTCTGCTTTAAAAAAT 3230  
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; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
; Polymorphisms in the Human Genome
; FILE REFERENCE: 108827.129
; CURRENT APPLICATION NUMBER: US/10/027,632
; PRIOR FILING DATE: 2002-04-30
; PRIOR FILING DATE: 2000-07-12
; PRIOR FILING DATE: 2000-04-20
; PRIOR FILING DATE: 2000-03-29
; PRIOR FILING DATE: 2000-02-24
; PRIOR FILING DATE: 1999-11-23
; PRIOR FILING DATE: 1999-09-28
; PRIOR FILING DATE: 1999-08-09
; NUMBER OF SEQ ID NOS: 325720
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 208140
; LENGTH: 637
; TYPE: DNA
; ORGANISM: Human
US-10-027-632-208140

Query Match      12.8%; Score 611.8; DB 5; Length 637;
Best Local Similarity 99.2%; Pred. No. 2.4e-144;
Matches 632; Conservative 3; Mismatches 0; Indels 2; Gaps 2;

Qy 3289 CATGGTGATCTGTGATTTTCAAGACCTTTAAATCCA-TTTTGAAGAATCAATTTTCATA 3347
Db 1 CATGGTGATCTGTGATTTTCAAGACCTTTAAATCCATTTTGAAGAATCAATTTTCATA 60

Qy 3348 TTTGCAATGGGTGCGCATGTGGAAGAGTGATATGCTTTTGTGCTGTAGCTTCAGAAAG 3407
Db 61 TTTGCAATGGGTGCGCATGTGGAAGAGTGATATGCTTTTGTGCTGTAGCTTCAGAAAG 120

Qy 3408 CACAGGAGGGAGAGCAATGTTGTTCA-GAGAAAGATCAACAGGAGGAGAACTGTCAGAG 3466
Db 121 CACAGGAGGGAGAGCAATGTTGTTCAAGAGATCAACAGGAGGAGAACTGTCAGAG 180

Qy 3467 CTGCTGAAATAGGGTGGTGGGAGGCATTAATCCCTCTCGTTGGGGGTAAAGCAG 3526
Db 181 CTGCTGAAATAGGGTGGTGGGAGGCATTAATCCCTCTCGTTGGGGGTAAAGCAG 240

Qy 3527 AACCGAGGTTGGTAGTAAATGTCATGACAGACAGTAGAGGGAGCAATAAATCTTAAATTTCT 3586
Db 241 AACCGAGGTTGGTAGTAAATGTCATGACAGACAGTAGAGGGAGCAATAAATCTTAAATTTCT 300

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Db 301 TTATAGTCTTGGAGTCTTTTGAGATAGAAAAGAAATATCTTTTGGCCCTTATGTCATAAAGAA 360

Qy 3647 GTATGGAAGGTGAAAGCGGAGAAAGCAGGAGAAAGCAATGATGATATATATATAGA 3706
Db 361 GTATGGAAGGTGAAAGCGGAGAAAGCAGGAGAAAGCAATGATGATATATATATAGA 420

Qy 3707 GGACAACTGTCAGAGGTTTTTCTTGAATATATGCAAAATATGATAGATTTAGGAAATTC 3766
Db 3766 GGACAACTGTCAGAGGTTTTTCTTGAATATATGCAAAATATGATAGATTTAGGAAATTC 480

Qy 3767 AGTAGGGAAATGCTTTTCACTTTGGGTTTTCTCTTCGATTAAGTTTGGGATCTCTCA 3826
Db 481 AGTAGGGAAATGCTTTTCACTTTGGGTTTTCTCTTCGATTAAGTTTGGGATCTCTCA 540

Qy 3827 TCTGCAATTTGACTTGGAGAGAGAAAGATGATGTTAGGACCTATATCTGTTTCTATT 3886
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US-10-027-632-208141
; Sequence 208141, Application US/10027632
; Publication No. US20020198371A1
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
; Polymorphisms in the Human Genome
; FILE REFERENCE: 108827.129
; CURRENT APPLICATION NUMBER: US/10/027,632
; CURRENT FILING DATE: 2002-04-30
; PRIOR FILING DATE: 2000-07-12
; PRIOR FILING DATE: 2000-04-20
; PRIOR FILING DATE: 2000-03-29
; PRIOR FILING DATE: 2000-02-24
; PRIOR FILING DATE: 1999-11-23
; PRIOR FILING DATE: 1999-09-28
; PRIOR FILING DATE: 1999-08-09
; NUMBER OF SEQ ID NOS: 325720
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 208141
; LENGTH: 637
; TYPE: DNA
; ORGANISM: Human
US-10-027-632-208141
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Query Match      12.8%; Score 611.8; DB 5; Length 637;
Best Local Similarity 99.2%; Pred. No. 2.4e-144;
Matches 632; Conservative 3; Mismatches 0; Indels 2; Gaps 2;

Qy 3289 CATGGTGATCTGTGATTTTCAAGACCTTTAAATCCA-TTTTGAAGAATCAATTTTCATA 3347
Db 1 CATGGTGATCTGTGATTTTCAAGACCTTTAAATCCATTTTGAAGAATCAATTTTCATA 60

Qy 3348 TTTGCAATGGGTGCGCATGTGGAAGAGTGATATGCTTTTGTGCTGTAGCTTCAGAAAG 3407
Db 61 TTTGCAATGGGTGCGCATGTGGAAGAGTGATATGCTTTTGTGCTGTAGCTTCAGAAAG 120

Qy 3408 CACAGGAGGGAGAGCAATGTTGTTCA-GAGAAAGATCAACAGGAGGAGAACTGTCAGAG 3466
Db 121 CACAGGAGGGAGAGCAATGTTGTTCAAGAGATCAACAGGAGGAGAACTGTCAGAG 180

Qy 3467 CTGCTGAAATAGGGTGGTGGGAGGCATTAATCCCTCTCGTTGGGGGTAAAGCAG 3526
Db 181 CTGCTGAAATAGGGTGGTGGGAGGCATTAATCCCTCTCGTTGGGGGTAAAGCAG 240

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Db 241 AACCGAGGTTGGTAGTAAATGTCATGACAGACAGTAGAGGGAGCAATAAATCTTAAATTTCT 300

Qy 3587 TTATAGTCTTGGAGTCTTTTGAGATAGAAAAGAAATATCTTTTGGCCCTTATGTCATAAAGAA 3646
Db 301 TTATAGTCTTGGAGTCTTTTGAGATAGAAAAGAAATATCTTTTGGCCCTTATGTCATAAAGAA 360

Qy 3647 GTATGGAAGGTGAAAGCGGAGAAAGCAGGAGAAAGCAATGATGATATATATATAGA 3706
Db 361 GTATGGAAGGTGAAAGCGGAGAAAGCAGGAGAAAGCAATGATGATATATATATAGA 420

Qy 3707 GGACAACTGTCAGAGGTTTTTCTTGAATATATGCAAAATATGATAGATTTAGGAAATTC 3766
Db 421 GGACAACTGTCAGAGGTTTTTCTTGAATATATGCAAAATATGATAGATTTAGGAAATTC 480
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Db	181	CTGTCTGAATAGGGTGGTTTGGAGGCAATTAATCCCTCTGTTGGGGTAAAGCAG	240
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Qy	3587	TTATAGTCTGGAGTCTTTGAGATAGAAAAGAAATATCTTTTGGCCCTTATGTCAAAAGAA	3646
Db	301	TTATAGTCTGGAGTCTTTGAGATAGAAAAGAAATATCTTTTGGCCCTTATGTCAAAAGAA	360
Qy	3647	GTATGGAAGAGTGAAGCGCGGAAGAACGACGAGGAGGAGAACCACTGATTAATATAGA	3706
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Qy	3707	GGACAATGGTGACAAGGTTTTCTTGAAATTAATGCAAAATATGATAGATTAGAGGAATTC	3766
Db	421	GGACAATGGTGACAAGGTTTTCTTGAAATTAATGCAAAATATGATAGATTAGAGGAATTC	480
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Db	481	AGTAGGGAATGCTTTTCACTTGAATTTGGGTTTCCCTCTCGATTAAAGTTTGGGATCCTCA	540
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Qy	3887	AACTAAAGCAAGTGGAAAAGACTTATTTGGTATTTTT	3923
Db	601	AACTAAAGCAAGTGGAAAAGACTTATTTGGTATTTTT	637

RESULT 15

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US-10-027-632-208141
; Sequence 208141, Application US/10027632
; Publication No. US20030204075A9
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
; FILE OF INVENTION: Polymorphisms in the Human Genome
; FILE REFERENCE: 108827.129
; CURRENT APPLICATION NUMBER: US/10/027,632
; CURRENT FILING DATE: 2002-04-30
; PRIOR APPLICATION NUMBER: US 60/218,006
; PRIOR FILING DATE: 2000-07-12
; PRIOR APPLICATION NUMBER: US 60/198,676
; PRIOR FILING DATE: 2000-04-20
; PRIOR APPLICATION NUMBER: US 60/193,483
; PRIOR FILING DATE: 2000-03-29
; PRIOR APPLICATION NUMBER: US 60/185,218
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: US 60/167,363
; PRIOR FILING DATE: 1999-11-23
; PRIOR APPLICATION NUMBER: US 60/156,358
; PRIOR FILING DATE: 1999-09-28
; PRIOR APPLICATION NUMBER: US 60/146,002
; PRIOR FILING DATE: 1999-08-09
; NUMBER OF SEQ ID NOS: 325720
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 208141
; LENGTH: 637
; TYPE: DNA
; ORGANISM: Human
US-10-027-632-208141

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Query Match	12.8%	Score 611.8;	DB 6;	Length 637;
Best Local Similarity	99.2%	Pred No. 2,4e-144;		
Matches 632;	Conservative 3;	Mismatches 0;	Indels 2;	Gaps 2;
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Db	1	CATGGTGATCTGTGTGATTTTCAAGACCTTTAATCCAATTTTGTGAAAGAAATCAATTTTCATA	60	

Search completed: December 21, 2005, 21:08:41  
Job time : 2425.07 secs

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Qy	3408	CACAGAGGGAGAGCAATGTTGTTCA- GAGAAAGATCAACAGAGGAGAACTGTCTAGAG	3466
Db	121	CACAGAGGGAGAGCAATGTTGTTCA- GAGAAAGATCAACAGAGGAGAACTGTCTAGAG	180
Qy	3467	CTGTCGAAATAGGGTGGTTTGGGAGGCAATTAATCCCTCTCGTTGGGGTAAAAAGCAG	3526
Db	181	CTGTCGAAATAGGGTGGTTTGGGAGGCAATTAATCCCTCTCGTTGGGGTAAAAAGCAG	240
Qy	3527	AACGACGGTGGTAGTAAATGCAATGACAGACAGTAGGGGACGATAAACTTTAAATTC	3586
Db	241	AACGACGGTGGTAGTAAATGCAATGACAGACAGTAGGGGACGATAAACTTTAAATTC	300
Qy	3587	TTATAGTCTGGAGTCTTTGAGATAGAAAAGAAATATCTTTTGGCCCTTATGTCAAAAGAA	3646
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Qy	3647	GTATGGAAGAGTGAAGCGCGGAAGAACGACGAGGAGGAGAACCACTGATTAATATAGA	3706
Db	361	GTATGGAAGAGTGAAGCGCGGAAGAACGACGAGGAGGAGAACCACTGATTAATATAGA	420
Qy	3707	GGACAATGGTGACAAGGTTTTCTTGAAATTAATGCAAAATATGATAGATTAGAGGAATTC	3766
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Db	481	AGTAGGGAATGCTTTTCACTTGAATTTGGGTTTCCCTCTCGATTAAAGTTTGGGATCCTCA	540
Qy	3827	TCTGCATTTGACTTTGGAGAGAGAAAGAAATGAATGTTAGGACCTATATCTGTTTCTATT	3886
Db	541	TCTGCATTTGACTTTGGAGAGAGAAAGAAATGAATGTTAGGACCTATATCTGTTTCTATT	600
Qy	3887	AACTAAAGCAAGTGGAAAAGACTTATTTGGTATTTTT	3923
Db	601	AACTAAAGCAAGTGGAAAAGACTTATTTGGTATTTTT	637



GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 334.473 Seconds  
(without alignments)  
7442.822 Million cell updates/sec

Title: US-09-751-797-25

Perfect score: 4797

Sequence: 1 tgcacaagcagaattcttcag.....gatgcccaagcgattttt 4797

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 4168288 seqs, 259477437 residues

Total number of hits satisfying chosen parameters: 8336576

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

- Published Applications NA New:\*
- 1: /cgn2\_6/ptodata/2/pubpna/US08\_NEW\_PUB.seq:\*
  - 2: /cgn2\_6/ptodata/2/pubpna/US06\_NEW\_PUB.seq:\*
  - 3: /cgn2\_6/ptodata/2/pubpna/US07\_NEW\_PUB.seq:\*
  - 4: /cgn2\_6/ptodata/2/pubpna/PCT\_NEW\_PUB.seq:\*
  - 5: /cgn2\_6/ptodata/2/pubpna/US09\_NEW\_PUB.seq:\*
  - 6: /cgn2\_6/ptodata/2/pubpna/US10\_NEW\_PUB.seq:\*
  - 7: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq:\*
  - 8: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq:\*
  - 9: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq:\*
  - 10: /cgn2\_6/ptodata/2/pubpna/US60\_NEW\_PUB.seq:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	4797	100.0	4797	7	US-11-177-987-26
2	686	14.3	7445	7	US-11-177-987-8
3	650	13.6	5935	7	US-11-177-987-42
4	258	5.4	690	7	US-11-177-987-25
5	159.4	3.3	418	7	US-11-177-987-18
6	244	5.1	1152	7	US-11-102-240-153
7	127.6	2.7	1111	7	US-11-177-987-9
8	126	2.6	1119	7	US-11-177-987-7
9	121.6	2.5	164527	7	US-11-121-086-71
10	118.6	2.5	130472	6	US-10-995-561-13312
11	114	2.4	180654	7	US-11-121-086-58
12	113.8	2.4	179777	7	US-11-121-086-106
13	112.4	2.3	191091	7	US-11-121-086-60
14	110.4	2.3	398287	6	US-10-995-561-13396
15	107.8	2.2	150450	7	US-11-112-908-54
16	107.8	2.2	191343	7	US-11-112-908-53
17	107.6	2.2	222094	6	US-10-995-561-13244
18	107.4	2.2	40394	6	US-10-995-561-13493
19	106.8	2.2	199321	7	US-11-121-086-10
20	106.4	2.2	169495	7	US-11-121-086-61
21	105.8	2.2	196200	7	US-11-121-086-9
22	104.8	2.2	105550	6	US-10-995-561-13235
23	103.6	2.2	1125000	6	US-10-995-561-13286

ALIGNMENTS

RESULT 1

US-11-177-987-26  
; Sequence 26, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; INTERLEUKIN-21, The Proteins Encoded, and Uses Thereof  
; FILE OF INVENTION: LUD 5664  
; CURRENT FILING DATE: 2005-07-07  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; PRIOR FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 26  
; LENGTH: 4797  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-11-177-987-26

Query Match 100.0%; Score 4797; DB 7; Length 4797;  
Best Local Similarity 100.0%; Pred. No. 0;  
Matches 4797; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db	121	CCACCAAGTCCCTCTTCTTCTTGGCCCTCTTGTGTACAGGAGGAGCAGCTGCGCCATCA 180
Qy	181	GTTCCCACTGACGCTTGACAACTTCAGAGCCCTATATACCAACCGCACT 240
Db	181	GTTCCCACTGACGCTTGACAACTTCAGAGCCCTATATACCAACCGCACT 240



Db 181 GCTCCCACTGAGGCTTGACAAGTCCAACTTCAGCAGCCCTATATCAACCAACCGCACCT 240  
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 Db 241 TCATGCTGGCTAAGGAGGTATACATCTCAATCTCTGCTCTTTCTCGTTGGATCTACTTTGGA 300  
 Qy 301 ATCCAAATAGTCTTAAACTTTCTTCAGAGCATCTCTAAGAGCTTTTAAAGAACCCACTGT 360  
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 Qy 361 TTATCCCTGAGGGTAGATAAAATTTCTGTTTTTTCAGAGACTCTTTGGGAATCTGGCTTT 420  
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 Db 481 CCCAAGAGCGGCATTCAGTAATCCATCTGATGATTTTTTTTTTCCCTTATGCTCTGTG 540  
 Qy 541 CATTTGTTCTAACTCATGCACACATCTGAATCTCTGCTTTTAGTCTTTATGATGTTGCTCT 600  
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 Db 601 GGGAGACGGATGGGGCACATGCTATGATATAAATTTTTTTTTTCTATTTGCTCAATGTCC 660  
 Qy 661 AGACCCCTAGTCTTTCTCTTCCAGGCTAGCTTGGCTGATACCAACACAGACGTTTCG 720  
 Db 661 AGACCCCTAGTCTTTCTCTTCCAGGCTAGCTTGGCTGATACCAACACAGACGTTTCG 720  
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 Db 721 TCTCATTTGGGAGAAACTGTTCCACGGAGTCAGTGTAAAGCTACAGTTGTGACGAACAGGG 780  
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 Db 1021 TCAGTCTTCAGCTATGCGACCTTACCCCTCTTCCCTTCCACAGACCCCTTAC 1080  
 Qy 1081 CCCAACTCTCTCTCTCCCTTACCCCTTACCCCTTACCCCTTACCCCTTACCCCTTAC 1140  
 Db 1081 CCCAACTCTCTCTCTCTCCCTTACCCCTTACCCCTTACCCCTTACCCCTTACCCCTTAC 1140  
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 Db 1381 ATTAGAAGAGAAAGTGGGAAATCGGAAAGGCTTAAAGTTCGGTGGTGGTCCGCGACTGTT 1440  
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 Qy 1501 CATTTGAAACTATGAAAAAAGTTTGAGTGCAGTGGGCCCCAGTAAAGAGGCCCTAGGACTT 1560  
 Db 1501 CATTTGAAACTATGAAAAAAGTTTGAGTGCAGTGGGCCCCAGTAAAGAGGCCCTAGGACTT 1560  
 Qy 1561 ACTGAAGAGGCTTAAATTTTTCATAGATGTTTATGTAATTTCTTGTCTTAAGCATG 1620  
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 Qy 1621 CAATTTTCTGAGATACGATTTGAGTTTATTTCTTACAGATTTGTCATAAATCTACTCCG 1680  
 Db 1621 CAATTTTCTGAGATACGATTTGAGTTTATTTCTTACAGATTTGTCATAAATCTACTCCG 1680  
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 Db 1681 CTCTTTCCACAAATGCAAACTCAGTAGGATTTCCCAAGATGAAGAGAGGTCTCTTGTA 1740  
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 Db 1741 AGGGAAGTGAAGTCTGAGTTCTGGCGTCCAAAGGAAATTCAGAGCTCAGGAAATCTAGGTCAC 1800  
 Qy 1801 TGTGAAATCTAGGTCATTTGTTGGGCAAAATTAATAAGAGCTTTAAATTTCCAGGTCGAAATGT 1860  
 Db 1801 TGTGAAATCTAGGTCATTTGTTGGGCAAAATTAATAAGAGCTTTAAATTTCCAGGTCGAAATGT 1860  
 Qy 1861 ACTGTACTCTCATGGGTGGAGGTTCTATAAGTTTCAGACACAAATTAAGATAGTTATG 1920  
 Db 1861 ACTGTACTCTCATGGGTGGAGGTTCTATAAGTTTCAGACACAAATTAAGATAGTTATG 1920  
 Qy 1921 CTTGTTATTTGTTTATAGCATATTAAGAGTGAACCTGCATATCCAGAGAAATGTGCAA 1980  
 Db 1921 CTTGTTATTTGTTTATAGCATATTAAGAGTGAACCTGCATATCCAGAGAAATGTGCAA 1980  
 Qy 1981 AAGCTGAAGGACAGTGAAGGAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGT 2040  
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 Qy 2041 AGGAGACAAATGTTGTTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTA 2100  
 Db 2041 AGGAGACAAATGTTGTTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTA 2100  
 Qy 2101 CTTGATTTCTCTACACAGGCGGATTAATTTGGTGTCTGTGTATGTAGATATATCTATA 2160  
 Db 2101 CTTGATTTCTCTACACAGGCGGATTAATTTGGTGTCTGTGTATGTAGATATATCTATA 2160  
 Qy 2161 TATCTAGATGTCAGTTTCCAAATCTTGCAAAATTTCTAGAAATTTCTAGAACTGGTGGGATCT 2220  
 Db 2161 TATCTAGATGTCAGTTTCCAAATCTTGCAAAATTTCTAGAAATTTCTAGAACTGGTGGGATCT 2220  
 Qy 2221 TAGCTTCTAGTCACATTAACCTCAGATTTCTGGGGATGCTCAGTGGCAGAGATAGGGCTA 2280  
 Db 2221 TAGCTTCTAGTCACATTAACCTCAGATTTCTGGGGATGCTCAGTGGCAGAGATAGGGCTA 2280  
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 Db 2281 GAATGCAAGTCTCTGTAATCCCAAGCCAGCAGCTTTTCCCGGTGGTGTATACAGATTTAGTTT 2340  
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 Db 2341 TGGTACCAATTTTCTTAGGGAAAAATTTTCCAGATTTCTTATTTGACTCATGTAACTGAGAGAG 2400

QY	2401	TACTTGTTTAAACACAGAAAATGCCCTATCGGCACAAATTTATTTGAAGTCATTTTGAAGT	2460
Db	2401	TACTTGTTTAAACACAGAAAATGCCCTATCGGCACAAATTTATTTGAAGTCATTTTGAAGT	2460
QY	2461	CATTAAATGCATGCTGTTGAAACTTGGAAAGTAATAACTCAGAAACAATGAGAAAAGAGCTGG	2520
Db	2461	CATTAAATGCATGCTTGTGAACTTGGAAAGTAATAACTCAGAAACAATGAGAAAAGAGCTGG	2520
QY	2521	ACTTGCCATATAGGGCTAAATTTCTGGAGTAATAAACAACCTTATTTTGAATTAATCATATATC	2580
Db	2521	ACTTGCCATATAGGGCTAAATTTCTGGAGTAATAAACAACCTTATTTTGAATTAATCATATATC	2580
QY	2581	TATCAGATATGATATAGTTTAAAGCAAGAGCAGACAACCCCGCATCTCTTTTATACAG	2640
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QY	2701	GGTAAGCTTTTTTCTCTCTCTCCCATCAAGACCTTCCATCTAGTATTTCTTCTTCA	2760
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Db	2761	CTCCCTCAACAAATCCCTAGGAGCATTTATCCATGGTGGCTGGTGTACATTTCTATAG	2820
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QY	2881	ACAATAGTGACTCACCCCAAAACCGGAGGAATGATTAGGAGCAGTGAAAGTGCAGCTCTT	2940
Db	2881	ACAATAGTGACTCACCCCAAAACCGGAGGAATGATTAGGAGCAGTGAAAGTGCAGCTCTT	2940
QY	2941	GCAAGCAGGTACAACTAAATCTCAGAAAACATGAAGCTCCAGTGTATGTAATTTTCAGT	3000
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Db	3001	AACAAGCTTAACTTAACTCCCTTTTCCCTCTTGACTTTTAAAGACGCTTCTTC	3060
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QY	3121	TTAAATGTGAAGCCCAAGTCTCTTGTTATAGAACTATTAATCTAGACATGGAGGGCTGAA	3180
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QY	3181	TGTTAGCATGCCACAGCAAGGCATGCTTTTACACATCTTGCTTAAAGAAATCTGATTC	3240
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Db	3241	ATCTTGCTGCTGTCTTTTGAAGTGAAGTGTGAGAGAGGAATCTCATGTGTGATCTG	3300
QY	3301	TGTGATTTTCAAGACCTTTTAAATCCATTTTGAAGAAATCAATTTTCAATTTTGAATGGGTT	3360
Db	3301	TGTGATTTTCAAGACCTTTTAAATCCATTTTGAAGAAATCAATTTTCAATTTTGAATGGGTT	3360
QY	3361	GCCATGTGGAAGATGATTAATGCTTTTCTGCTGGTAGCTTCAGAAAGCAGAGGGAGA	3420
Db	3361	GCCATGTGGAAGATGATTAATGCTTTTCTGCTGGTAGCTTCAGAAAGCAGAGGGAGA	3420
QY	3421	GCAATGTTGTTTACAGAAAAGATCAACAGGAGGAAACTGTCTAGAGCTGTCTGAAATAGG	3480
Db	3421	GCAATGTTGTTTACAGAAAAGATCAACAGGAGGAAACTGTCTAGAGCTGTCTGAAATAGG	3480
QY	3481	GTGGTTTTGGAGGCATTAATTTCCCTCTCGTTGGGGTAAAGCAGAACGCAGGTTGGTA	3540

Db	3481	GTGGTTTTGGAGGCATTAATTTCCCTCTCGTTGGGGTAAAGCAGAACGCAGGTTGGTA	3540
QY	3541	GTAATAATGCATGACACACAGTAGTAGGGACGATAAACTTTAAAAATCTTTATAGTCTTGGAG	3600
Db	3541	GTAATAATGCATGACACACAGTAGTAGGGACGATAAACTTTAAAAATCTTTATAGTCTTGGAG	3600
QY	3601	TCCTTGAGATAGAAAAGATATCTTTTGGCCCTTATGTCAAAGAGATATGAAAAGGTGA	3660
Db	3601	TCCTTGAGATAGAAAAGATATCTTTTGGCCCTTATGTCAAAGAGATATGAAAAGGTGA	3660
QY	3661	AAGGGCGGAGAAAGCAGGAAAGGAGAACCATGTATTATATAGAGGACAATGGTGACA	3720
Db	3661	AAGGGCGGAGAAAGCAGGAAAGGAGAACCATGTATTATATAGAGGACAATGGTGACA	3720
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Db	3721	AGGTTTTTCTTGAAAATAATGCAAAATATGATAGAGGAAATTCAGTAGGGAATGCTT	3780
QY	3781	TTCACTTGAATTTGGGTTTCTCTTCGATTAAGTTTGGGATCTCATCTGCAATTTGACTT	3840
Db	3781	TTCACTTGAATTTGGGTTTCTCTTCGATTAAGTTTGGGATCTCATCTGCAATTTGACTT	3840
QY	3841	GGAGAGAAAAGATGAATGTTAGGACCTATATCTGTTTTCTATTAACTAAAGCAAGTG	3900
Db	3841	GGAGAGAAAAGATGAATGTTAGGACCTATATCTGTTTTCTATTAACTAAAGCAAGTG	3900
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Db	3961	AAAAGTGAAAATAGAAAAGCCTTAATGTATTTGGTGAATACATGTTTCAAAGTCATTTG	4020
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QY	4081	GAAATCTCTTGGTACACATAAAATACAAAGAACATAGCTGATGAGCTAAAAAGTCCA	4140
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QY	4141	TGCATAAATCTCATACTGTTTTTAAAGAAAGTTTATGAATTTCTGTAGGGTGCAATTCAAAG	4200
Db	4141	TGCATAAATCTCATACTGTTTTTAAAGAAAGTTTATGAATTTCTGTAGGGTGCAATTCAAAG	4200
QY	4201	CTGTCTCGGCCCATGTGCGGCCCTGTGGGCTGCAAGTTTGGACAAGCTCTTTATAGTAAATC	4260
Db	4201	CTGTCTCGGCCCATGTGCGGCCCTGTGGGCTGCAAGTTTGGACAAGCTCTTTATAGTAAATC	4260
QY	4261	TGTCATAGATAGTTTTTGGAGCTGCAAAAACAGGCCCAAGGCATATGTTGGTGGCCTCGGGAT	4320
Db	4261	TGTCATAGATAGTTTTTGGAGCTGCAAAAACAGGCCCAAGGCATATGTTGGTGGCCTCGGGAT	4320
QY	4321	CCCCAGATCCCAAGCCTCACTTCAGTCTCTCTGCTCTGTTTAAAGAGGGTGCTCAACTC	4380
Db	4321	CCCCAGATCCCAAGCCTCACTTCAGTCTCTCTGCTCTGTTTAAAGAGGGTGCTCAACTC	4380
QY	4381	TCGTGCCAGCTTTTAAACAGCTTCATTAGTGTAGGTGCAACCTGAAATTTGATGCCCTGCTG	4440
Db	4381	TCGTGCCAGCTTTTAAACAGCTTCATTAGTGTAGGTGCAACCTGAAATTTGATGCCCTGCTG	4440
QY	4441	GTGGCCTCTCAGTCCAGAGAGCGCTCATTTTAAAGCTCTTTGGCAAAATCATACAATACTAA	4500
Db	4441	GTGGCCTCTCAGTCCAGAGAGCGCTCATTTTAAAGCTCTTTGGCAAAATCATACAATACTAA	4500
QY	4501	AGGGATTTTACTATGAATGTTTTTACAAATGCTTTAAAGCTCGGTTTCTGTCTCCATCAACC	4560
Db	4501	AGGGATTTTACTATGAATGTTTTTACAAATGCTTTAAAGCTCGGTTTCTGTCTCCATCAACC	4560
QY	4561	TAACTTGTCAATTTCTTAAATTTGTCTCTTTAGAAAACATGGGCAATAAATGCTTCAAACTTT	4620



Qy	1461	AGCCACAAAATCGGAGCGGTGTGAACCTTGATGCGCGTGAAACATTTGTAAGAACTATGAAAAA	1520
Db	3478	AGTTAAGAAAAACCAAGT-GTGTGAGTTTGATGTCTTCAGACACACCCCAACTATGAAACAT	3536
Qy	1521	AGTTTGAGTGGAGTGGGCCCCAGTAAAGGCCCTAGGACTTACTTGAAGAGGGCTTAATTTT	1580
Db	3537	ATCCAGGAGGACGGCGAGACTGTGGAGACCTTGGCAATTTTAGGGA--AGCGCGGGCTTTT	3594
Qy	1581	CACATGAGATGTTTTATGTGACATTTCTTGTTCTAAGCATGCAATTTTTCTGGAGATACGAT	1640
Db	3595	CACAGAGAACTTTATGCTCATCTCTTGCTACATCTCCACCTTTGATGAGGTTTCAGC	3654
Qy	1641	TGAGGTTTTATTCCTTACAGAAATTTGCATAAACTACCTCCGCTCTTCCCAAAATGCAAAC	1700
Db	3655	TCAGGTTTCGTTTTCT-----ACCGTTCTTGCTACTGTTGGGAAC	3693
Qy	1701	CTCAGTAGGATTTCCCAAGATGAAGAGAGGTCTCTTTGTAAGGGAAGTGATCGATTTCTG	1760
Db	3694	TTTCAGTAGGATTTCCCAAGACGAGGACAGCTCTTCTGTGAAGGAGGACCTCGATTTTCA	3753
Qy	1761	GCGTCCAAGGGAATTCAGAGAGCTCAGGAATCTAGGTCACTGTTTGAATCTAGTCAATG	1820
Db	3754	GTGTCTTAGAGAACGAATAGCTCAGGAATCTAGGTCAAGTGAATCTAGGTCAACAGC	3813
Qy	1821	TGGGCAAAATTTACTAAGAGCTTTAAATTCACGGTGAAATGTACTGTACTCCATCGGCTG	1880
Db	3814	GGGCAAAATGCACTGAACGCTCTTAATTCAGGTGAACGGTCACGTGCTCAGATATAC	3873
Qy	1881	GAGGTTTCATAAAGTTTCAGCAACAATTAAGATAGTTATGCTGTTATGTTTATATAGCA	1940
Db	3874	AGSTATTGGGCTCCACCGGATAAGATCTGTTAGTGA-GTCTGCTTTTATTTTTCAGCA	3932
Qy	1941	TATTGAAGTGTACACCTGCATATCCAGAGGAATGTCAAAGCTGAAGACACACGTGA	2000
Db	3933	CATCAGCGGTGACACACAGAAATCTCAGAGAATGTGAGAAAGGCTGAAGGAGACAGTGA	3992
Qy	2001	AAAGGTAGGACTGATAACTGTCAATGCTCAAGTCAATGCAATAGGAGAGACAAATGTTG	2060
Db	3993	AAAGTACTATTGGCAAGCCACATACTAAGCCATTCACTAGTAG-GAGACGTGGGATTTC	4050
Qy	2061	TTCTTTCCTTTCTTTCTCCCATCACTTTGTGATTTTTTCACTTGATTTCTCTACCAACCAG	2120
Db	4051	TTTCTCTGCTTCCAGTCCCTTCTACTTTGTGAACATTTTATTTGACTTGTCTACTATCTG	4110
Qy	2121	GGCGATTA----CTTTGGTGTGTGTATGTAGATATATCTATATATCTAGATGTCAGTT	2176
Db	4111	GTCCATTTACTCGCTTAGCTGCACCTGTATCTAGCTGGGTCTATAGATCTTTCAATCTGTG	4170
Qy	2177	TCCAAATCTTGCAAAATGTAGAAATCTAGAACTGGTTGGGATCTTAGCTTGTCTAGTCAAC	2236
Db	4171	TCATAATTT---GTAAGTCAAAATCTGAGAGTACGAGAAAGCTTAGTCAGCAGTCTC	4227
Qy	2237	ATAACCTCAGATTTCTGGGATGGTCACTGGCAGAGATAGGGCTAGAAATGAGGTCCTCTG	2296
Db	4228	ATGAGCACTTGCTCGGAGGATGGCTTGTGACAGAGTCAATGCTAGAGAAGCAGCATCCCTG	4287
Qy	2297	AATCCCAAGCCAGACATTTTCCCGGTGGTGAATACAGATTAGTTTTTGTGATCAATTAATCT	2356
Db	4288	ATTCCCAAGCTCTGCAC-TTGCTAGTGGCCATGTGTAAATTACTTTGGCTTGATTAAGTAT	4346
Qy	2357	TAGGGAATTTTCAGATTCCTATTGACTCATGTATCTGAAGAGTACTCTGTTTAAABAACA	2416
Db	4347	TTGGGAAA--GCCAGTTTCCACGACCTCATATAATCTGAAGAACCATGCAATGAAACATA	4404
Qy	2417	GAAAAATGCCCTATGGGCAAAATTTATTTGAAGTCAATTTTGAAGTCAATTAATGCAATGCTT	2476
Db	4405	GAAA-----GCTGGGCAAACTTACTAGAGATGATTTTGGCTCATTAACCGATGCTC	4460
Qy	2477	TGAAACTTGGAGAATAAACTCAGAAACAATGAGAAAAAGCTGGACTTGCATATAGGGCT	2536
Db	4461	TGAAATGTGGCAAAATCAACCCAGAAATAACAACAAAAGAGCTGGATTTGCAATAGGACA	4520

QY	2537	AA	TTCTCGGA-----GTAATAAACACCTATTATTTTGAATTTATCATTAATATCTATCAGATA	2589
DB	4521	AG	TATTTAGAAATCACCTGGTATTAATAGCTATCATCTTAATTTAAATATATAGGGCTATATA	4580
QY	2590	TT	GATTATAGTTTAAAGCAAGACGACACAACC-CGAGTCTCTTTTATACAGGTTTCAAAT	2648
DB	4581	TAT	TATTTAAGATTAAACACAAGAGTGGATAGCTTCCCAATTATCTTGGCCTGGTTTTCAA	4640
QY	2649	AG	AGTAAAAAATATTAGTAAAGAGATTTATTATAGTTAAATGGAAGTCTGAAATTCGTAACT	2708
DB	4641	AG	AGTAAAAAATATCAGTCATGGATTAATATATAGTGTCTCATGAAAGTATGAGATGGAACACC	4700
QY	2709	TT	TTTTTCTTCCTCTCTCCCATCAGACCTTCATCTAGTTTCTTCCTTCACTCCCTCA	2768
DB	4701	TT	TCCTTACTTTTTTACCTCA-----TTTCTTAGTTTTTTTTTTCTTTCAACCCCTGA	4752
QY	2769	AC	AAATCCCTAGGAGCATTTATCCATGCTGGCTGTGTACATTTCTATAGTGAATGAT	2828
DB	4753	TCA	AGCCACTAGTAAGCACTATCTGCTGTGAGCTATTATATGACTTTACGMAAACAC	4812
QY	2829	ACC	ATCATATGTGGCTATTTTGGTGAAAGAAACA--ACAATGGAAGGCTTAGACTAACAAATA	2886
DB	4813	ATT	GCTGTGTGGCTCTCTTCGSGAAGGGAACAGATAGCAGGAGGCTCAGCTAGCAAGT	4872
QY	2887	GT	ACTCACCCCAAAACCGGAGGAATGATTAGAGACGATGAAGTGAAGCTCTT-GCAAG	2945
DB	4873	CT	GACTTGGCCTTAAAGCCAGAGGCGATGGTTGATAGCAGAGAAAGTGAAGCTCTTCGCAAG	4932
QY	2946	CAG	GTACAACTAATATCTCAGAACATGAAGGCTCCAGTTGATGGAAATTTTCAGTAACAA	3005
DB	4933	TGG	GTGTCTTAAGTAAATCAGAAACAGGAAGGCTCCGGTTGATGGAAATATCATAGTAAGAT	4992
QY	3006	GC	TTAACTTAAATTCGCCCTTTTTTCCCTCTCTGACTTTTTTAAAAAAGCGTTTCTTCCTGAG	3065
DB	4993	AT	CTACCTCTTATCTCCTTCTATCGAAC-----TAAATCGTCTCTTTCTTTCTGTG	5042
QY	3066	CAT	CAITTAATAGATGTGACTGTTTCTCTCTTGAATAATTGAAGGCTTTGTAGTTTTAAA	3125
DB	5043	TG	TAGGCTGATAAACACACTTGT--TTCTTTTGAGTGTTCATGGCTTTGTAGATTTTAA	5100
QY	3126	TT	TGAAGCCAGTCTCTCTGTTTATAGAACTATTATCTAGACATGGAAGGCTGAAATGTTA	3185
DB	5101	GT	CTCTGCCAGTCTTGTGTAGAG--GGTTGTGTACCTTGACACCTGGGCTTGATGTTA	5158
QY	3186	GC	ATGCCACAGACAAGGCATGCTTTACACATCTTGCTTAAAAAATTTACTGATTTCACTTT	3245
DB	5159	GC	ATGCCAAAGGCACACACTCTGATGCTCTGTGTAAAGGTTATTATTATTACT--	5215
QY	3246	CT	TGTTGTCTTTAGAAAAGTGAAGTGTGAGAGAGGAATCTCATGGTGA-----	3296
DB	5216	----	TTGCTTTTGGAAAGGTGAAGCGTGTGTGAGAAAGAACTCACAGGAGATGTGTCT	5270
QY	3297	-----	TCGTGTGATTTTTCAGACCTTTATCCACTTTTATCCATTTTGAAGATCAATT	3342
DB	5271	CT	GTAGAAAACCTTTTTTTTTTCCCTTTAAATGCTCTATAATCCACTTTTCAGTCAAC--	5327
QY	3343	TC	ATTTTGCATGGGTTGTCATGTGGAAGATGATTATGCTTTTTTGTGCTGAGTTCA	3402
DB	5328	TG	ACTTTTATACCATGCTGTGTCACTGMAAGAGTGTTTAGGCCGCTCTCATGCTCTGGG	5387
QY	3403	GAA	AGCA-CAGGAGGGAGACAATGTTGTTTCAGAGAAAGATCAACAGGAGGAGAAACTGT	3461
DB	5388	AAA	AGCAACCAATAGGGGAAGGAATGTTATGCTCAGAAATCTGACCGGACGGGAAACTGGT	5447
QY	3462	CAG	AGCTGTCTGAAATAGGGTGGTTTTTGGAGGCAATTAAATCCCTCTCGTTGGGGTAAA	3521
DB	5448	CAG	AGCTCTCCCCGAAGACCA-----CCACAGGTGTTTAAGTAGG	5485
QY	3522	AG	CAGAAGCAGGTTGGTAGTAAAT--GCATGACAGACAGTACGGGACGATAAATTTAA	3580
DB	5486	AAC	AGTCCAGGTTGGGCTCAATGTAATAGAAATGAAACGACGAGGGAAGATATAGCTACAA	5545
QY	3581	AA	TTCTTTTATAGTCTTGGAGTCTTTGAGATAGAAAAGAAATATCTTTTTTGGCCTTATGTCA	3640



Db 764 CTTGGCCTTCAGGACACATATCTGAATTTTATCTACAGAGGCCGTTT--AGAAAGCCA 821  
Qy 509 CTGATGATTTTTTTTCCCTTATGCTCTGTCGCAATTTCTAAACTCATGACACATCTG 568  
Db 822 CCCACGACTGCAATACCTTCCATCCTGTTGCTCTCTCTGAACCTATCTCTTTGGC 881  
Qy 569 AATTCTGCTTTTGTCTTTATGATGTTGCTCTGGGAGACGGGATGGGACACATGCTAT 628  
Db 882 TACTC-----CTGAGACCCACTGCGGACATACATCTCTAC 916  
Qy 629 GTATAAATTTTTTTTCTATTGCTCAATGTCACACCTTAGTCTTTTCTTCTCTCCAG 688  
Db 917 TTACAGGCTTTTCTTCATCTCTCTGTCACCCAGGCACTTAGGGTTTTC-TCTCTTTTCA 975  
Qy 689 GCTAGCTTGGCTGATAACAACACAGAGGTTGCTCTCAATGGGGAGAACTGTTCCACGGA 748  
Db 976 GCCAGCCTTCAGATAACAACACAGAGCTCCGGCTCATCGGGAGAACTGTTCCGAGGA 1035  
Qy 749 GTCAAGTGAAGTACAGTTGTGACGAACAGAGCCGCTGTGCGCTCCATGGGTACTTGGGT 808  
Db 1036 GTCAAGTGAAGTCTCACTGTGATGAGCAGGC-----TAGCTGCGGAGCT 1082  
Qy 809 GGTGCTGATGATGTTTAGGCTTTATCCCTTATGACCCCTTCTGTTTCCCTTCCACCTGC 868  
Db 1083 GGTGACCCCTCTGGATAG----TCTGACGTATGACCCCTGCTGCTCTTGTCTACCTGC 1138  
Qy 869 AGATGAGTGAAGCGCTGCTATCTGATGAAGCAGGTGCTGAACCTCACCCCTTGAAGAGTGC 928  
Db 1139 AGGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCACTTCACTCCCTGGAGACATC 1198  
Qy 929 TGTTCCTCAATCTGATAGGTTTCCAGCCTTATATGACAGAGGTGGTGGCTTCTCGGCCA 988  
Db 1199 TGTCTCCCACTCAGACAGGTTTCCGGCCCTACATGACAGAGGTGGTGGCTTCTCGACCA 1258  
Qy 989 GGCTCAGCAGCAGCTAAGCAGATGCTAAGTTGAGTCTCAGCCTATGCCCTACCTACCC 1048  
Db 1259 AACTCAGCAATCAGCTCAGCTCTGTGTAGTCTGGCTCTGGCTACCTATGCTCTCTCT 1318  
Qy 1049 CTCCTCCCTCTTCCACAGAGCCCTTACCACCACTCTCTCTCTCTCTCCCTCCCTACCCC 1108  
Db 1319 CTTCTCTTCTATTCAGTAGAACCCGAGTCTCTGCTCTCTCTCTCTCTCTCTCTCTCTCT 1378  
Qy 1109 TAAGCTAGCAGGAAGAAGTGTCTTGGCAGCAGTGTATCAGGAGTCA-----TTTGGG 1161  
Db 1379 GGAGGGCTCAGCACCAACCACATATAGGCCACTTGAAATAGGTCAAAAGGCTTTGGC 1438  
Qy 1162 ATCATAGATATTGCTTTGCTTTGCTGACTGAGTCACATCTGAGTTTATAGTGTGAATG 1221  
Db 1439 TTCAATTGAGTAATCTTGAATTTGATTATGATTAGCTTTATTTGTTTATCATCGAA 1498  
Qy 1222 GGGTCTGGAACCTTAAGTGTACAGAGCCGCAATGGTTGTTCTCGGAAAAAAGGCAACTC 1281  
Db 1499 AGAATCACTCAATTTCTGTAGATGAGAAGATGTGGGACAAAAGGCTAGAT 1558  
Qy 1282 AGGTGCGTAA--GATGAGAAAGGTGTGGGAAAAATCTAGCTGTGGAATGGATCCA 1338  
Db 1559 AGAAAAACAGATCTGCTGAGTACAGTACTTATGGGGGGGGGGGAGGCGGATATCCA 1618  
Qy 1339 TTGAGTCTAAGTTGTTGAGGGGGGGGATGGCTGAGAGAAATAGAGAGAAAGTGG 1398  
Db 1619 CTGAGTCAAGTACTTGTGGGAGAGAAATCCACTGAGTACAAGTACTTGTGGGGGAAG 1678  
Qy 1399 AAATGGGAAGCTTAAAGTGGTGGTGGGCGAGACTGTGCCC-----TGTTGA 1450  
Db 1679 AATGGCACAGCAAAAGTTGAAGGGAAAGAGGAAGATGAGAGGCCCTCAATGTTGGGG 1738  
Qy 1451 TGTCAATGGGAAGCCAAAAATCGAGCGGTGTGAATTTGATGCGCTGGAACATTTGAAAC 1510  
Db 1739 TGTAAAGGTCACCTCTTTTCCATGTGTATGGAGAGTTAAGAAAAATCAGTGTGTGAGTT 1798  
Qy 1511 TATGAAAAAAGTTTGTAGTGGAGTGGGCCCACTGTAAGAGCCCTAGGACTTACTGAAGG 1570  
Db 1799 TGATGTCTTCAGACACCCCACTATGCGAGCTGTGGGAGACCTGGGCAATTTAGGGA-AGG 1857

Qy 1571 GCTTAATTTTTCATGAGATGTTTTTATGTACATTTCTTGTCTTAAGCATGCAATTTTCTG 1630  
Db 1858 CGGGCTTTTTCACAGAGAAATTTATGCTCATCTCTGTGTCTACACTCCACCTTTGAT 1917  
Qy 1631 GAGATAGATTTGAGTTTATTTCTTACAGAAATTTGATAACTACTCCGCTCTTTCCAC 1690  
Db 1918 GAGTTAAGCTCAGTTTCTGTTCT-----ACCGTTCTTGTGTAC 1956  
Qy 1691 AAATGCAAACTCAGTAGGATTTCCCAAGATGAAGAGAGTCTCTTGTAAAGGGAAGTGA 1750  
Db 1957 TGGTGGAACTTTCAGTAGGATTTCCCAAGACGAGGACAGCTCTTCTGTAAAGGAGGAC 2016  
Qy 1751 CTGATTTCTGCGCTCCAAAGGGAATTTCAAGAGCTCAGAAATTTAGGTCACTGTTGAAATC 1810  
Db 2017 CTGATTTCTAGTCTCTAGAGAACGAAATAGCTCAGAGAAATCTAGGTCAACGCTGAATCT 2076  
Qy 1811 TAGGTCAATTTGGGCAAAATTTACTAGAGCTTTAAATCCAGGTGAATTTGACTTACTCTC 1870  
Db 2077 AGTCAAGCGGGCAAAATTTGACTGAACGCTCTTATCCAGGTGAACGGTCCACGTGCTC 2136  
Qy 1871 CATGGGTGTGAGGTTTCAAAAGTTTTCAGCAACAATTAAGATAGTTATGCTGTATTG 1930  
Db 2137 AGATATACTAGGATTTGGGCTCCACCGGATAGATTTCTGTAGTGA-GTCTGCTTTTA 2195  
Qy 1931 TTTTATAGCATATTTGAAGGTGATGACCTGTCATATCCAGAGGAATTTGCAAAAGCTGAAG 1990  
Db 2196 TTTTGCAGCACATCAGTGGTGAACGACCAGAAATCCAGAAAGATGTCAAGAGGCTGAAG 2255  
Qy 1991 ACACAGTGAAGAGTGAAGTGAATGATGATCTCAATGCTAGTCAATGCAATAGGAGAGACA 2050  
Db 2256 AGACAGTGAAGAGTGAATGATGATGATGATGATGATGATGATGATGATGATGATGATG 2315  
Qy 2051 AATGTTGTTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2110  
Db 2316 GGGATTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2373  
Qy 2111 CTACCAAGGCGGATTT-----ACTTTGGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 2166  
Db 2374 CTACTGTCTGGTCCATTTACTCACTTAGCTGCACCTGCATCTAGCTGGGTCTATAGATCTT 2433  
Qy 2167 GATGTCAGTTTCCAAATCTTGCAAATTTGTAGAAATCTAGAACTGGTGGGATCTTAGCTT 2226  
Db 2434 TCAATCTGTCTTAAATTT---GTAAGTCAAAATTTCTGGAGCTAGCAGAAAGCTTAGCTC 2490  
Qy 2227 GTCTAGTCAATAACCTCAGATTTCTGGGATGTTCTGTCGAGAGATAGGCTAGAGATGC 2286  
Db 2491 AGCAGTCTCTGAGCACTTGTCTGGAGGATGGCTTGTGACAGAGTCAATGCTAGAGAC 2550  
Qy 2287 AGGTCTCTGAATCCCAAGCCAGCACCTTTTCCGGTGGTGATACAGATTTAGTTTGGTAC 2346  
Db 2551 AGCATCCCTGATTTCCAGCTCTGCAC--TTGCCCTAGTGGCCACGTTAATTTAGCTT 2609  
Qy 2347 CATTAATTTTAGGAAATTTTCAATTTCTTATGATCTATGATCTATGATCTGAGAGAGTCT 2406  
Db 2610 GATTAAGTATTTGGGAAA--GCCAAATTTCCACCCACCTACATATCCGAAAGAGCTAGCA 2667  
Qy 2407 TTTAAAAACAGAAAAATGCTCTATGGGCAATTTTATTTGAAAGTCAATTTTGAAGTCATTAA 2466  
Db 2668 TTGAAAACTAGAAA----GCTGGGCAAAAATTTACTAGAGATGATTTTGGAGTCAATTA 2723  
Qy 2467 TGCATTTGCTTTGAACTTGAAGAATAAATCTAGAAACAATGAGAAAAAGAGCTGAGCTGC 2526  
Db 2724 ACTGATGCTCTGAAAATGTGATCAATCAACCCAGNATAACAACAAAGAGAGCTGAGTTGC 2783  
Qy 2527 ATATAGGCTAAATTTCTGGAGTAAATAACACTTAT-----TTTGAATTTATCATATA 2578  
Db 2784 AAATAGGACAAAGTATTTAGAAATCACTGGTATTAACAGCTGTCTCATTTTAAATAATAG 2843  
Qy 2579 TCT---ATCAGATATTGATTTATAGTTTAAAGCAAGAGAGCAGACAAAC--CCGATCTCTTT 2634  
Db 2844 TGCTATTTAGTCCCTTATTTAAGATTTAAACAAAGAGTGGATTAATTTCCCAATTTTACTG 2903

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Qy 2635 ATACAGGTTCAAATAGAGTAAATAATTAGTAGAGATTTATTATTAGTTAAATGGAAGTC 2694
Db 2904 GGCCTGGTTTCAATAGAGTAAATAATATACGTATAGATTAATATAGTGTGTCATGAAAGTA 2963
Qy 2695 TGAATTGGTAAGCTTTTTTTTTTCTCTCTCTCCCATCAAGACCTTCCATTCTAGTTTCTT 2754
Db 2964 TGAGTTGGAACCC---CTTTCTTACTTTTACCTTTCATTCTTAGTTATTATTTTTTTT 3020
Qy 2755 CTTTCACTCCCTCAACAAATCCCTAGGAGATTTATCCATGGTGGGCTGGTGTACATTT 2814
Db 3021 TCTTCAACCCCTGATCAAGCCACTAGTAAGCACCTATCTGTGTGAGCTATFATATGACT 3080
Qy 2815 CTATAGTCAATGATACCATCATGCGCTATTCTGTGAAGAACA--ACAATGGAAGGC 2872
Db 3081 TTACAGCAAAACAATTCGTGTGGCTCTTTTGGGAAGGGAACAGATAGCAGGAGGC 3140
Qy 2873 TTAGACTAACAA--TAGTGACTCACCCCAAAACCGGAGGAATGATTAGGAGCAGTGAAAGT 2931
Db 3141 TCAGGCTAGCAAGTCTGGACTCAACCTAAAGCCAGAGGCGATGGTTGATAGCAGAGAAAGT 3200
Qy 2932 GACGCTCTT--GCAAGCAGGTACAATAATACTCAGAAACATGAAGGCTCAGTTGATGG 2990
Db 3201 GAGGCTCTTCAACAAGTGGGTGTCTTAAGTAATCAGAAACAGGAAGGCTCTGGTTGATGG 3260
Qy 2991 AATTTTCAGTAAACAAGCTTAACCTTAATTCCTCTCTTCCCTTTTTCGACTTTTTTAAAAA 3050
Db 3261 AATTATCAGTAAGATATCTACCTTAICTCC-----TTCCTTATAGAAGCTTAACCG 3313
Qy 3051 GCGTTTCTCTCTGAGCATCAATTAATAGTGTGACTGTTCTCTCTTTGTAATGAAGG 3110
Db 3314 TCTCTCTCTCTGTGTGAGCTGATAAACAGCGTTGTT--TTCCTTGAAGTTCATGG 3371
Qy 3111 CTTTGTAGTTTAAATGTAAGCCAGTCTCTTGTGTATAGAATAATATCTAGACATG 3170
Db 3372 CTTTGCAGATTTTCAGTCTCTGCGAGTCTTGT--TAGAGGGTTTGTACTTTGCACACC 3429
Qy 3171 GAGGCGTAAATGTAGCATGCCACAGCAAGGCATGCTTTACACATCTTGCTTAAAAAT 3230
Db 3430 TGGGCTTGGATGTAGCATGCCAAAGGCACACACTTCTGAATGCTGTGTAAAAGTTAT 3489
Qy 3231 TACTGATTTTCATCTTGTCTTGTGTTTGTCTTTAGAAAAAGTGAAGTGTGAGAGAGGAATCTCA 3290
Db 3490 TATTCATTTACT-----TTGCTTTGGAAAGTGAAGTGTGTGTGAGAAAGAACTCA 3541

RESULT 4
US-11-177-987-25
; Sequence 25, Application US/11177987
; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 25
; LENGTH: 690
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-11-177-987-25
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Query Match 5.4%; Score 258; DB 7; Length 690;
Best Local Similarity 100.0%; Pred. No. 7.1e-59;
Matches 258; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TGACAAAGCAGAGATCTTTTCAGAACAGAGTTCTCTTCCCAAGTACCAGTTCAGTTAG 60
Db 1 TGACAAAGCAGAGATCTTTTCAGAACAGAGTTCTCTTCCCAAGTACCAGTTCAGTTAG 60
Qy 61 AATTGTCTGCAANTGGCGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGG 120
Db 61 AATTGTCTGCAANTGGCGCCCTGCAGAAATCTGTGAGCTCTTTCTTATGGGACCCCTGG 120
Qy 121 CCACCAAGTGCCTCTCTCTTCTTGGCCCTCTTGGTACAGGAGGAGCAGCTGCGCCCATCA 180
Db 121 CCACCAAGTGCCTCTCTCTTCTTGGCCCTCTTGGTACAGGAGGAGCAGCTGCGCCCATCA 180
Qy 181 GCTCCCACTGAGGCTTTGACAAAGTCCAACCTTCCAGCAGCCCTATATCACCACCGCACCT 240
Db 181 GCTCCCACTGAGGCTTTGACAAAGTCCAACCTTCCAGCAGCCCTATATCACCACCGCACCT 240
Qy 241 TCATGCTGGCTAAGGAGG 258
Db 241 TCATGCTGGCTAAGGAGG 258

RESULT 5
US-11-102-240-153
; Sequence 153, Application US/11102240
; Publication No. US20050260647A1
; GENERAL INFORMATION:
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin I.
; APPLICANT: Wood, William I.
; TITLE OF INVENTION: ANTIBODIES TO POLYPEPTIDES ENCODED BY A NUCLEIC ACID UNDEREXPRESS
; TITLE OF INVENTION: ESOPHAGEAL TUMOR
; FILE REFERENCE: P3230R1C106C
; CURRENT APPLICATION NUMBER: US/11/102,240
; CURRENT FILING DATE: 2005-04-08
; PRIOR APPLICATION NUMBER: 10/063662
; PRIOR FILING DATE: 2002-05-07
; PRIOR APPLICATION NUMBER: 10/006867
; PRIOR FILING DATE: 2001-12-06
; PRIOR APPLICATION NUMBER: PCT/US00/23328
; PRIOR FILING DATE: 2000-08-24
; PRIOR APPLICATION NUMBER: 60/170262
; PRIOR FILING DATE: 199-12-09
; NUMBER OF SEQ ID NOS: 170
; SEQ ID NO 153
; LENGTH: 1152
; TYPE: DNA
; ORGANISM: Homo Sapien
US-11-102-240-153

Query Match 5.1%; Score 244; DB 7; Length 1152;
Best Local Similarity 100.0%; Pred. No. 5.5e-59;
Matches 244; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 15 CTTCAGAACAGGTTCTCTTCCCAAGTCCACAGTTGCTCGAGTTAGAAATTGTTGCAATG 74
Db 1 CTTCAGAACAGGTTCTCTTCCCAAGTCCACAGTTGCTCGAGTTAGAAATTGTTGCAATG 60
Qy 75 GCCGCCCTGCAGAAATCTGTGAGCTCTTCTTATGGGACCCCTGGCCACCCAGCTGCCTC 134
Db 61 GCCGCCCTGCAGAAATCTGTGAGCTCTTCTTATGGGACCCCTGGCCACCCAGCTGCCTC 120
Qy 135 CTTCTCTTGGCCCTCTTGGTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGCAAG 194
Db 121 CTTCTCTTGGCCCTCTTGGTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGCAAG 180
Qy 195 CTTGACAGTCCAACTTCCAGCAGCCCTATATCACCACCGACCTTATGCTGCTAAG 254
Db 195 CTTGACAGTCCAACTTCCAGCAGCCCTATATCACCACCGACCTTATGCTGCTAAG 254
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Db 181 CTTGACAGTCCAACTTCCAGCAGCCCTATATACCAACCGCACCTTCATGCTGGCTAAG 240
Qy 255 GAGG 258
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Db 241 GAGG 244

RESULT 6
US-11-177-987-18
; Sequence 18, Application US/11177987
; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 18
; LENGTH: 418
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-11-177-987-18

Query Match 3.3%; Score 159.4; DB 7; Length 418;
Best Local Similarity 99.4%; Pred. No. 1.1e-32;
Matches 160; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 4637 ACAGCTTGGAGAGTGGAGAGATCAAGCAATTTGGAGAACTGGATTGCTGTTTATGTC 4696
Db 157 AAAGCTTGGAGAGTGGAGAGATCAAGCAATTTGGAGAACTGGATTGCTGTTTATGTC 216

Qy 4697 TCTGAGAAATGCTGCAATTTGACAGAGCAAGCTGAAAAATGAATACTAACCCCTTT 4756
Db 217 TCTGAGAAATGCTGCAATTTGACAGAGCAAGCTGAAAAATGAATACTAACCCCTTT 276

Qy 4757 CCTGCTAGAAATAACAATTAGATGCCCAAGCGATTTT 4797
Db 277 CCTGCTAGAAATAACAATTAGATGCCCAAGCGATTTT 317

RESULT 7
US-11-177-987-9
; Sequence 9, Application US/11177987
; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43

US-11-177-987-9

Query Match 2.6%; Score 126; DB 7; Length 1119;
Best Local Similarity 71.%; Pred. No. 1.9e-23;
Matches 165; Conservative 0; Mismatches 65; Indels 0; Gaps 0;

Qy 29 CTCCTTCCCAGTCACAGTTGCTCGAGTTAGAAATGCTCGCAATGGCGCCCTGCAGAA 88
Db 9 CTCCTCTCTCACTATCAACTGTTGACACTTGTGCGATCTCTGATGGCTCTCTGCAGAA 68

Qy 89 ATCTGTGAGCTCTTTCTTATGGGACCCCTGATGGCCACAGCTGCTCTCTCTTTGGCCCT 148
Db 69 ATCTATGAGTTTTCCTTATGGGACTTTGGCCGACGCTGCTCTCTCTCTCTCTCTCT 128

Qy 149 CTTGGTACAGGGAGGAGCAGCTCGGCCCATCAGCTCCCACTGAGCTTGCAGAGTCCAA 208
Db 129 GTGGGCCCAAGGAGGCAAAATGCGCTGCCCGTCAACACCCGGTGAAGTGAAGTGTCAA 188

Qy 209 CTTCCAGCAGCCCTATATCAACCAACCGCACCTTTCATGCTGGCTAAGGAGG 258
```

```
; SEQ ID NO 9
; LENGTH: 1111
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-11-177-987-9

Query Match 2.7%; Score 127.6; DB 7; Length 1111;
Best Local Similarity 72.3%; Pred. No. 7e-24;
Matches 166; Conservative 0; Mismatches 64; Indels 0; Gaps 0;

Qy 29 CTCCTTCCCAGTCACAGTTGCTCGAGTTAGAAATGCTCGCAATGGCGCCCTGCAGAA 88
Db 7 CTCCTCTCTCACTATCAACTTGTGACACTTGTGCGATCGGTGATGGCTGTCTCTGCAGAA 66

Qy 89 ATCTGTGAGCTCTTTCTTATGGGACCCCTGATGGCCACAGCTGCTCTCTCTTTGGCCCT 148
Db 67 ATCTATGAGTTTTCCTTATGGGACTTTGGCCGACGCTGCTCTCTCTCTCTCTCTCT 126

Qy 149 CTTGGTACAGGGAGGAGCAGCTCGGCCCATCAGCTCCCACTGAGCTTGCAGAGTCCAA 208
Db 127 GTGGGCCCAAGGAGGCAAAATGCGCTGCCCATCAACACCCGGTGAAGTGTCAA 186

Qy 209 CTTCCAGCAGCCCTATATCAACCAACCGCACCTTTCATGCTGGCTAAGGAGG 258
Db 187 CTTCCAGCAGCCGTACATCGTCAACCGCACCTTTCATGCTGGCCCAAGGAGG 236

RESULT 8
US-11-177-987-7
; Sequence 7, Application US/11177987
; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 7
; LENGTH: 1119
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-11-177-987-7

Query Match 2.6%; Score 126; DB 7; Length 1119;
Best Local Similarity 71.%; Pred. No. 1.9e-23;
Matches 165; Conservative 0; Mismatches 65; Indels 0; Gaps 0;

Qy 29 CTCCTTCCCAGTCACAGTTGCTCGAGTTAGAAATGCTCGCAATGGCGCCCTGCAGAA 88
Db 9 CTCCTCTCTCACTATCAACTGTTGACACTTGTGCGATCTCTGATGGCTCTCTGCAGAA 68

Qy 89 ATCTGTGAGCTCTTTCTTATGGGACCCCTGATGGCCACAGCTGCTCTCTCTTTGGCCCT 148
Db 69 ATCTATGAGTTTTCCTTATGGGACTTTGGCCGACGCTGCTCTCTCTCTCTCTCTCT 128

Qy 149 CTTGGTACAGGGAGGAGCAGCTCGGCCCATCAGCTCCCACTGAGCTTGCAGAGTCCAA 208
Db 129 GTGGGCCCAAGGAGGCAAAATGCGCTGCCCGTCAACACCCGGTGAAGTGAAGTGTCAA 188

Qy 209 CTTCCAGCAGCCCTATATCAACCAACCGCACCTTTCATGCTGGCTAAGGAGG 258
```



Db 189 CTTCCAGCAGCGTACATGTCACCGCACCTTTATGCTGGCCAAAGGAG 238  
|||||

## RESULT 9

US-11-121-086-71/c  
; Sequence 71, Application US/11121086  
; Publication No. US20050266459A1  
; GENERAL INFORMATION:  
; APPLICANT: POULSEN, TIM S.  
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES  
; FILE REFERENCE: 09138.6000-00000  
; CURRENT APPLICATION NUMBER: US/11/121,086  
; CURRENT FILING DATE: 2005-05-04  
; PRIOR FILING DATE: 2004-05-04  
; NUMBER OF SEQ ID NOS: 107  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 71  
; LENGTH: 164527  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-11-121-086-71

Query Match 2.5%; Score 121.6; DB 7; Length 164527;  
Best Local Similarity 75.8%; Pred. No. 7.9e-21; Mismatches 49; Indels 8; Gaps 2;  
Matches 179; Conservative 0;  
Qy 4027 GATGTTTAAATCAGGAGTGTCCTCATTTGGCTTCCCTGGACACCTTGA-----AA 4080  
Db 116406 GGTGTTCTAAATAGGGGTGTCGATCTTTTCGGCTTCCCTGGCCACATTGGAAGAAGAA 116347  
Qy 4081 GAATGTCTTGGTACACATAAATAACAAG--ACATAGCTGATGAGCTTAAAGATC 4138  
Db 116346 GAATGTCTTGGCCACACATAAATACTTAACACATAGCTGATGAGCTTAAAGAAAT 116287  
Qy 4139 CATGCATTAATCTCATCTGTTTAAAGAAAGTTTATGAAATTTCTGTTAGGCTGATTCAA 4198  
Db 116286 CACAAAAAATCTCTTAATGTTTAAAGAACTTTACAAATTTGTTGGCCGCGATTCAA 116227  
Qy 4199 AGCTGTCTGGCCATGTCGGCTGTGGGCTGTCAGAGTTGGACAGCTTCTTTATAA 4254  
Db 116226 AGCGTCTTAGCCACATGTGGCCATGGCCATGGGTTGCAAGTTGTTCTTAA 116171  
|||||

## RESULT 10

US-10-995-561-13312  
; Sequence 13312, Application US/10995561  
; Publication No. US20050272054A1  
; GENERAL INFORMATION:  
; APPLICANT: CARGILL, Michele et al.  
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH  
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF  
; TITLE OF INVENTION: DETECTION AND USES THEREOF  
; FILE REFERENCE: CL001559  
; CURRENT APPLICATION NUMBER: US/10/995,561  
; CURRENT FILING DATE: 2004-11-24  
; NUMBER OF SEQ ID NOS: 85702  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 13312  
; LENGTH: 130472  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-10-995-561-13312

Query Match 2.5%; Score 118.6; DB 6; Length 130472;  
Best Local Similarity 74.3%; Pred. No. 4.3e-20; Mismatches 54; Indels 8; Gaps 2;  
Matches 179; Conservative 0;  
Qy 4031 TTTTAAATCAGGAGTGTCCTCATTTGGCTTCCCTGGACACCTTGAAGAAGATTTGCTT 4090  
Db 11104 TTGTAGATCAGGGGTGTCCTTCTTTGGCTTCCCGGCCACATTGGAAGAATTGTCTT 11163  
|||||

Qy 4091 GGTACACACATAAAAT-----ACAAGAACAAATAGCTGATGAGC--TAAAAAGTCATG 4142  
Db 11164 GGGCCACAGATAAAATACACTAACATAGCTGATAAGCTTTAAAAAAACTGCAA 11223  
Qy 4143 CATAAATCTCATACTGTTTAAAGAAAGTTTATGAATTTCTGTTAGGCTGATTCAGAGCT 4202  
Db 11224 AAAAAATCTCACAGTGTGTTTAAACAAAGTTTATGAATTTGTTGGGTCACCTTTCAAAGCT 11283  
Qy 4203 GTCCTGGCCATGTGCGGCTGTGGCTGCAGGTTGGACAAGCTCCTTATAAGTAATCTG 4262  
Db 11284 GTGGTGGCTGCAATGAGCCCTAGGCTGTGGGTTGGACAAGCTTCTGTAGATTTATT 11343  
Qy 4263 T 4263  
Db 11344 T 11344  
|||||

## RESULT 11

US-11-121-086-58/c  
; Sequence 58, Application US/11121086  
; Publication No. US20050266459A1  
; GENERAL INFORMATION:  
; APPLICANT: POULSEN, TIM S.  
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES  
; FILE REFERENCE: 09138.6000-00000  
; CURRENT APPLICATION NUMBER: US/11/121,086  
; CURRENT FILING DATE: 2005-05-04  
; PRIOR FILING DATE: 2004-05-04  
; NUMBER OF SEQ ID NOS: 107  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 58  
; LENGTH: 180654  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-11-121-086-58

Query Match 2.4%; Score 114; DB 7; Length 180654;  
Best Local Similarity 75.8%; Pred. No. 9e-19; Mismatches 45; Indels 14; Gaps 3;  
Matches 185; Conservative 0;  
Qy 4028 ATGTTTTAAATCAGGAGTGTCCTCATTTGGCTTCCCTGGACACCTTGA-----AAG 4081  
Db 110206 ATATTTAGAACAGGTTGTCCTTCTTTGGCTTCCCTGGCCACATTGGAAGAAG 110147  
Qy 4082 AATTGCTCTGTGACACACATAAATAACAAGACA-----ATAGCTGATGAGCT--AAAA 4133  
Db 110146 AATTGCTCTGGGACACACATAAATAACAATACTAACGATTCGTGATGAGCTTAAAAA 110087  
|||||

Qy 4134 AAGTCCATGCATAAATCTCATCTGTTTAAAGAAAGTTTATGAATTTCTGTTAGGCTGCA 4193  
Db 110086 AAATCACAATAAAGTCTCTTAATGTTTAAAGAAAGTTTATGAATTTCTGTTAGGCTGCA 110027  
Qy 4194 TTCAAAGCTGTCTGGCCATGTGGGCTGTGGGCTGTGGGCTGTGGCAAGCTTCTTATA 4253  
Db 110026 TTCAAAGCTTCTTGGGCTGTGATGCAGCCCATGGTTCACGGGTTGGACAAGCTTATTAG 109967  
Qy 4254 AGTA 4257  
Db 109966 AGTA 109963  
|||||

## RESULT 12

US-11-121-086-106/c  
; Sequence 106, Application US/11121086  
; Publication No. US20050266459A1  
; GENERAL INFORMATION:  
; APPLICANT: POULSEN, TIM S.  
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES  
; FILE REFERENCE: 09138.6000-00000

; CURRENT APPLICATION NUMBER: US/11/121.086  
; CURRENT FILING DATE: 2005-05-04  
; PRIOR APPLICATION NUMBER: 60/567,570  
; PRIOR FILING DATE: 2004-05-04  
; NUMBER OF SEQ ID NOS: 107  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 106  
; LENGTH: 179777  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-11-121-086-106

Query Match 2.4%; Score 113.8; DB 7; Length 179777;  
Best Local Similarity 74.1%; Pred. No. 1e-18;  
Matches 177; Conservative 0; Mismatches 47; Indels 15; Gaps 2;  
Qy 4023 TAGAGATGTTTAAATCAGGAGTGTCCTCAATCATTTGGCTTCCCTGGACCACTTGAAGA 4082  
Db 120599 TAGAGATTTAGCTTCAGGACTGTTCAATCTTTTGGCTTCCCTGGCCACATTTGAAGA 120540  
Qy 4083 ATGCTCTGTGACACATATAAATACAGAACCA-----ATAGCTGATGAGCTAAAAAG 4136  
Db 120539 ATTGCTTGGTCACACATATAAATACAGTAACACTAATGATAGCTGTTGAATTAATAA 120480  
Qy 4137 TCCATGC-----ATAAATCTCATCTGTTTAAAGAAAGTTTATGAATTTCTGTAG 4187  
Db 120479 AAAAATCGCCAAAAAATTTTCAATATGTTTAAAGAAAGTTTATGAATTTGTGTGG 120420  
Qy 4188 GGTGCATTCAAAGCTGTCCTGGGCAATGTCGGGCTGTGGGCTGACAGTTGGACAGCT 4246  
Db 120419 GCCGCAATCAAGCCGCTTGGGCTGTCATGTGGCCCAAGGCGGCTGGGTTGAACAAGCT 120361

RESULT 13  
US-11-121-086-60  
; Sequence 60, Application US/11/121.086  
; Publication No. US20050266459A1  
; GENERAL INFORMATION:  
; APPLICANT: POULSEN, TIM S.  
; APPLICANT: NIELSEN, KIRSTEN V.  
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES  
; FILE REFERENCE: 09138.6000-00000  
; CURRENT APPLICATION NUMBER: US/11/121.086  
; CURRENT FILING DATE: 2005-05-04  
; PRIOR APPLICATION NUMBER: 60/567,570  
; PRIOR FILING DATE: 2004-05-04  
; NUMBER OF SEQ ID NOS: 107  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 60  
; LENGTH: 191091  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-11-121-086-60

Query Match 2.3%; Score 112.4; DB 7; Length 191091;  
Best Local Similarity 76.5%; Pred. No. 2.5e-18;  
Matches 179; Conservative 0; Mismatches 46; Indels 9; Gaps 3;  
Qy 4021 AGTAGAGATGTTTAAATCAGGAGTGTCCTCAATCATTTGGCTTCCCTGGACCACTTG--- 4077  
Db 147239 ATTATCCATGCCCTAGACAGGGGTGCCCATTTCTTTGGCTTCCCTGGGCTCTTTGGAA 147298  
Qy 4078 AAAGAATGTCTTGGTACACACATATAAATACAGAACCA-----ATAGCTGATGAGCTAAA 4132  
Db 147299 GAAGAATGTCTTGGTGCACATATAAATGCATTAACATTAACATAGCTAATAGCTAAA 147358  
Qy 4133 AAAGTCATGATATAATCTCATCTGTTTAAAGAAAGTTTATGAATTTCTGTAGGCTGC 4192  
Db 147359 AAAATCCCCCA-AAAATCTCATAGTGTGTTGAAGAAAGTTTACGAAATTTGTGTTGGTGC 147417  
Qy 4193 ATTCAAGCTGTCTCTGGGCAATGTCGGGCTGTGGGCTGACAGTTGGACAGCT 4246  
Db 147418 ATTCAAGCCATCTCTGGGCGGCATGAGGCTGTGGGCGGTGGGTTGGACAGGCT 147471

RESULT 14

US-10-995-561-13396  
; Sequence 13396, Application US/10995561  
; Publication No. US20050272054A1  
; GENERAL INFORMATION:  
; APPLICANT: CARGILL, Michele et al.  
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH  
; TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF  
; TITLE OF INVENTION: DETECTION AND USES THEREOF  
; FILE REFERENCE: CL001559  
; CURRENT APPLICATION NUMBER: US/10/995,561  
; CURRENT FILING DATE: 2004-11-24  
; NUMBER OF SEQ ID NOS: 85702  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 13396  
; LENGTH: 398287  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; NAME/KEY: misc feature  
; LOCATION: (1)-(398287)  
; OTHER INFORMATION: n = A,T,C or G, or insertion/deletion polymorphism (see Tables 1-  
US-10-995-561-13396

Query Match 2.3%; Score 110.4; DB 6; Length 398287;  
Best Local Similarity 73.3%; Pred. No. 1.4e-17;  
Matches 173; Conservative 0; Mismatches 51; Indels 12; Gaps 2;  
Qy 4036 AATCAGAGTGTCCTCAATCATTTGGCTTCCCTGGACCACTTG---AAAGAAATTTGCTTGG 4092  
Db 78384 AAGCTGGGTGTCCAAATCTTTTGGCTTCCCTGGGTCACTGGAAGAAGAAATCGCCTTGG 78443  
Qy 4093 TACACACATAAATACAGAACATAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCT 4152  
Db 78444 GCCACACATGAATAATATACTAACTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCT 78503  
Qy 4153 ATACTGT-----TTTAAGAAAGTTTATGAATTTCTGTAGGGTGCATTTCAAGCTG 4203  
Db 78504 AGAAATCTCATATATGTTTAAAGAAAGTTTACGAATTTGTTGGCGCGCTTTCAAGGCCA 78563  
Qy 4204 TCCTGGGCCATGTGCGGCTGTGCGGCTGCGGCTGCGGCTGCGGCTGCGGCTGCGGCTGCGGCT 4259  
Db 78564 CCCTGGGCTCATGCGGCGGTGGCCGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCT 78619

RESULT 15

US-11-112-908-54/c  
; Sequence 54, Application US/11112908  
; Publication No. US20050260659A1  
; GENERAL INFORMATION:  
; APPLICANT: Harris, Cole  
; APPLICANT: Davis, Lisa M.  
; TITLE OF INVENTION: Breast Cancer Biomarkers  
; FILE REFERENCE: 04-164-US  
; CURRENT APPLICATION NUMBER: US/11/112,908  
; CURRENT FILING DATE: 2005-04-22  
; PRIOR APPLICATION NUMBER: US 60/564,758  
; PRIOR FILING DATE: 2004-04-23  
; PRIOR APPLICATION NUMBER: US 60/575,978  
; PRIOR FILING DATE: 2004-06-01  
; PRIOR APPLICATION NUMBER: US 60/631,702  
; PRIOR FILING DATE: 2004-11-30  
; PRIOR APPLICATION NUMBER: US 60/633,826  
; PRIOR FILING DATE: 2004-12-07  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 54  
; LENGTH: 150450  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-11-112-908-54

Query Match	2.2%;	Score 107.8;	DB 7;	Length 150450;
Best Local Similarity	72.2%;	Pred. No. 3.6e-17;		
Matches 174;	Conservative 0;	Mismatches 52;	Indels 15;	Gaps 2;
QY	4037	ATCAGGAGTGTCCAATCATTTGGCTTCCCTGGACCACTTGA---AAGAATTGTCTTGGT	4093	
Db	101528	ATCAGGGGTGTCCAATCTTTTGGCTTCCCTGAGCCACATTGGGAAGAAGATTGTCTTGGG	101469	
QY	4094	ACACACATAAAATACAAGACA-----ATAGCTGATGAGCTAAAAAGTCCAT	4141	
Db	101468	CTACATATAACATACACTAACCTAATACTAATGATAGCTGATGAACCTTAAACAAAAAA	101409	
QY	4142	GCATAAATCTCATACTGTTTTAAGAAAAGTTTATGAATTTCTGTTAGGGTGCATTCAAAGC	4201	
Db	101408	GCAAAAAGTCTTAATGTTTTAGAAAAGTTTACAAAATTTGTGTGGTCCCAATTCAAAGC	101349	
QY	4202	TGTCCTGGGCCATGTGGCGGCTGTGGGCTGCAGGTGGACAAGCTCCTTATTAAGTAATCT	4261	
Db	101348	TATCTTGGACCACATGCAGCCTGTGGGTGTGAGGATAGACAAGCTTGCTTTAGATGATTT	101289	
QY	4262	G 4262		
Db	101288	G 101288		

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Job time : 355.473 secs

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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 671.791 Seconds  
(without alignments)  
15704.028 Million cell updates/sec

Title: US-09-751-797-29

Perfect score: 5935

Sequence: 1 gaattcaagtcacatgcaaa.....atattatctggtaaagttag 5935

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 1303057 seqs, 888780828 residues

Total number of hits satisfying chosen parameters: 2606114

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Issued Patents NA:\*

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- 2: /cgn2\_6/ptodata/1/ina/5 COMB.seq.\*
- 3: /cgn2\_6/ptodata/1/ina/6A COMB.seq.\*
- 4: /cgn2\_6/ptodata/1/ina/6B COMB.seq.\*
- 5: /cgn2\_6/ptodata/1/ina/H COMB.seq.\*
- 6: /cgn2\_6/ptodata/1/ina/PCTUS COMB.seq.\*
- 7: /cgn2\_6/ptodata/1/ina/PP COMB.seq.\*
- 8: /cgn2\_6/ptodata/1/ina/RE COMB.seq.\*
- 9: /cgn2\_6/ptodata/1/ina/backfiles1.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	5935	100.0	5935	3	US-09-178-973B-17
2	5935	100.0	5935	3	US-09-419-568F-29
3	5935	100.0	5935	3	US-09-354-243B-29
4	4245.2	71.5	7445	3	US-09-178-973B-8
5	4245.2	71.5	7445	3	US-09-419-568F-8
6	4245.2	71.5	7445	3	US-09-354-243B-8
7	651.8	11.0	8888	3	US-09-949-016-17185
8	650	11.0	4797	3	US-09-419-568F-25
9	650	11.0	4797	3	US-09-354-243B-25
10	602.4	10.1	1111	3	US-09-178-973B-9
11	602.4	10.1	1111	3	US-09-419-568F-9
12	602.4	10.1	1111	3	US-09-354-243B-9
13	555.2	9.4	1119	3	US-09-178-973B-7
14	555.2	9.4	1119	3	US-09-419-568F-7
15	555.2	9.4	1119	3	US-09-354-243B-7
16	555.2	9.4	1166	3	US-10-084-298-3
17	541.4	9.1	1050	3	US-10-090-365-40
18	218.4	3.7	1191	3	US-10-084-298-1
19	216.4	3.6	1116	3	US-10-090-365-14
20	216.4	3.6	1116	3	US-09-728-911-14
21	215	3.6	1152	3	US-09-870-574-1
22	171.8	2.9	601	3	US-09-949-016-190092
C 23	162	2.7	264665	3	US-09-949-016-13747
C 24	158.8	2.7	7218	2	US-08-232-463-14

c	25	154.2	2.6	2674	3	US-09-019-095A-1	Sequence 1, Appli
	26	153.4	2.6	270	3	US-09-443-282B-32	Sequence 32, Appl
	27	152	2.6	254	3	US-09-443-282B-33	Sequence 33, Appl
	28	151.8	2.6	1583	3	US-09-270-767-11302	Sequence 11302, A
	29	151.2	2.5	4598	2	US-07-807-043B-5	Sequence 5, Appli
	30	151.2	2.5	4598	2	US-08-299-849B-5	Sequence 5, Appli
	31	151.2	2.5	4698	2	US-08-142-368A-5	Sequence 5, Appli
	32	151.2	2.5	4698	3	US-08-967-727-5	Sequence 5, Appli
	33	151.2	2.5	4698	3	US-08-037-230D-5	Sequence 5, Appli
	34	151.2	2.5	4698	3	US-09-583-850-5	Sequence 5, Appli
	35	151.2	2.5	4698	3	US-09-579-197-5	Sequence 5, Appli
	36	151.2	2.5	4698	3	US-09-404-026-5	Sequence 5, Appli
	37	151.2	2.5	4698	3	US-09-312-464-5	Sequence 5, Appli
	38	151.2	2.5	4698	3	US-09-583-848A-5	Sequence 6, Appli
	39	140.2	2.4	2147	3	US-08-927-165A-6	Sequence 3, Appli
	40	137.2	2.3	3892	2	US-08-555-723B-3	Sequence 3, Appli
	41	137.2	2.3	3892	3	US-09-123-465-3	Sequence 1, Appli
	42	134.8	2.3	2509	3	US-09-319-284-1	Sequence 96, Appl
	43	133.6	2.2	7874	3	US-09-780-175-96	Sequence 96, Appl
	44	131.6	2.2	30310	3	US-09-657-346A-96	Sequence 5, Appli
	45	131.6	2.2	90050	3	US-09-245-041-5	Sequence 5, Appli

ALIGNMENTS

RESULT 1

US-09-178-973B-17  
; Sequence 17, Application US/09178973B  
; Patent No. 6274710

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; APPLICANT: Louhed, Jamila

; APPLICANT: Renauld, Jean-Christophe

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; TITLE OF INVENTION: (TIPS)

; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5543

; CURRENT APPLICATION NUMBER: US/09/178,973B

; CURRENT FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 17

; SEQ ID NO 17

; LENGTH: 5935

; TYPE: DNA

; ORGANISM: Mus musculus

US-09-178-973B-17

Query Match	100.0%;	Score	5935;	DB	3;	Length	5935;
Best Local Similarity	100.0%;	Pred. No.	0;				
Matches	5935;	Conservative	0;	Mismatches	0;	Indels	0;
Qy	1	GAATTCAGTCCACATGCAATCGAATACCTTTGTAATCTCTCTTCAAAATCC	60				
Db	1	GAATTCAGTCCACATGCAATCGAATACCTTTGTAATCTCTCTTCAAAATCC	60				
Qy	61	ATCTATATAGTATAAGTATTGATGATCATTTAAAAATAATGTTTGAGACTTATGTTT	120				
Db	61	ATCTATATAGTATAAGTATTGATGATCATTTAAAAATAATGTTTGAGACTTATGTTT	120				
Qy	121	GCAACAGTAAATGTCAGAGAGAAATGACAAATGATATAGTATTTATTTTAAAAAAT	180				
Db	121	GCAACAGTAAATGTCAGAGAGAAATGACAAATGATATAGTATTTATTTTAAAAAAT	180				
Qy	181	CTATGCTTAAATGCTTATAGTATTTCTACTACTGACATTTCCAACTTAACCTGACCT	240				
Db	181	CTATGCTTAAATGCTTATAGTATTTCTACTACTGACATTTCCAACTTAACCTGACCT	240				
Qy	241	TGCTATGATTTTCAACCTTTGATTTTGCACTACTACCACTGCTGCTCACTTACCATGC	300				
Db	241	TGCTATGATTTTCAACCTTTGATTTTGCACTACTACCACTGCTGCTCACTTACCATGC	300				
Qy	301	TATCCGACGACGATGTTCCCTCGATGTTTTCGCTTTTTCGCTCTCTCGCTAACGGCTCTC	360				

Db 301 TATCCGACGACGATGTTCCCTCGATGTTTGTGCTCTCTGCTCTCTCGCTAACAGGCTCTC 360  
Qy 361 CTCTCAGTTATCAACTTTTGACACTCTGCGATCGGTGATGCGTCTCTCGCAGAAATCTTA 420  
Db 361 CTCTCAGTTATCAACTTTTGACACTCTGCGATCGGTGATGCGTCTCTCGCAGAAATCTTA 420  
Qy 421 TGAGTTTTTCCCTTATGCGGACTTTGGCCGACGCTGCTCTCTCTCATTTGCCCTGTGGG 480  
Db 421 TGAGTTTTTCCCTTATGCGGACTTTGGCCGACGCTGCTCTCTCTCATTTGCCCTGTGGG 480  
Qy 481 CCCAGGAGCAAAATGCGCTGCCATCAACACCCCGGTGCAAGCTTGAGGTGCAAACTTCC 540  
Db 481 CCCAGGAGCAAAATGCGCTGCCATCAACACCCCGGTGCAAGCTTGAGGTGCAAACTTCC 540  
Qy 541 AGCAGCGTACATCGTCAACCGGACCTTTATGCTGGCCCAAGGAGGTACAGCTGCATCTCT 600  
Db 541 AGCAGCGTACATCGTCAACCGGACCTTTATGCTGGCCCAAGGAGGTACAGCTGCATCTCT 600  
Qy 601 TTCTCTCAVACCGCTTGGCCATTTCTGGAAGCACTTGCAAACTCTTTAGGGGCGCTTT 660  
Db 601 TTCTCTCAVACCGCTTGGCCATTTCTGGAAGCACTTGCAAACTCTTTAGGGGCGCTTT 660  
Qy 661 ATCTCCGAGGTCTCACTACCTATGTTTTCTGTCTCTTTAGAGACTCTTTAAGGACTGGA 720  
Db 661 ATCTCCGAGGTCTCACTACCTATGTTTTCTGTCTCTTTAGAGACTCTTTAAGGACTGGA 720  
Qy 721 TCTTTTCTATTTCTATTTCAAGTCTCAGGACCAATTTCTATCTGTCCTTTAGGACTGGA 780  
Db 721 TCTTTTCTATTTCTATTTCAAGTCTCAGGACCAATTTCTATCTGTCCTTTAGGACTGGA 780  
Qy 781 ATATACTGAATTTTATCTACAGAGCGGTTTTAGAAAGCCACCCACGACTGCAATATCTT 840  
Db 781 ATATACTGAATTTTATCTACAGAGCGGTTTTAGAAAGCCACCCACGACTGCAATATCTT 840  
Qy 841 CCATCTGTGTGCTCTCTCTGAACTCATACTCTCTTGCTACTCTCTGAGACCACTGC 900  
Db 841 CCATCTGTGTGCTCTCTCTGAACTCATACTCTCTTGCTACTCTCTGAGACCACTGC 900  
Qy 901 GGACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTCTGTCACCCAGGCACTTAGG 960  
Db 901 GGACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTCTGTCACCCAGGCACTTAGG 960  
Qy 961 TTTTCTCTCTTTTACGGCCAGCTTGCAGATAACAAACACAGACGCTCCGCTCATCGGGAG 1020  
Db 961 TTTTCTCTCTTTTACGGCCAGCTTGCAGATAACAAACACAGACGCTCCGCTCATCGGGAG 1020  
Qy 1021 AAACCTGTTCCGAGGAGTCAGTGAAGTCTCTCACTGTGATGAGCAGGCTAGCTCGGGAG 1080  
Db 1021 AAACCTGTTCCGAGGAGTCAGTGAAGTCTCTCACTGTGATGAGCAGGCTAGCTCGGGAG 1080  
Qy 1081 CTGTTGGACCTCTGGGATAGTCTGACGATAGACCCCTGCTGCTTCTGTCACCTGAG 1140  
Db 1081 CTGTTGGACCTCTGGGATAGTCTGACGATAGACCCCTGCTGCTTCTGTCACCTGAG 1140  
Qy 1141 GCTAAGGATCAGTCTCTCTCATGAAGCAGGTGCTCAACTTCACTTCCGGAAGACATCTG 1200  
Db 1141 GCTAAGGATCAGTCTCTCTCATGAAGCAGGTGCTCAACTTCACTTCCGGAAGACATCTG 1200  
Qy 1201 CTCCTCCAGTCAGACAGGTTCCGGCCCTTACATGAGGAGGTGGTGCCTTTCTGACCAAA 1260  
Db 1201 CTCCTCCAGTCAGACAGGTTCCGGCCCTTACATGAGGAGGTGGTGCCTTTCTGACCAAA 1260  
Qy 1261 CTCAGCAATCAGTCTCTGTTGAAGTCTGGCTCTGGCTACCTATGCTCTCTCTCT 1320  
Db 1261 CTCAGCAATCAGTCTCTGTTGAAGTCTGGCTCTGGCTACCTATGCTCTCTCTCT 1320  
Qy 1321 TCCTCTTCTATTTCCAGTAAGAACCGAGTCTGCTCTCTCTCTTCCAAAGAGTGAGG 1380  
Db 1321 TCCTCTTCTATTTCCAGTAAGAACCGAGTCTGCTCTCTCTCTCTTCCAAAGAGTGAGG 1380  
Qy 1381 AGGGCTCAGCACCAACCATCATAGGCCACTTGAATAGGTCAAAAGGCTTTGGCTT 1440  
Db 1381 AGGGCTCAGCACCAACCATCATAGGCCACTTGAATAGGTCAAAAGGCTTTGGCTT 1440

Qy 1441 CAATTGAGTAATACTTTGAGTTGTATAGTTAAGCTTTATTTGTTTTTATCCATGGAAG 1500  
Db 1441 CAATTGAGTAATACTTTGAGTTGTATAGTTAAGCTTTATTTGTTTTTATCCATGGAAG 1500  
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Db 1501 AAATCAACTCAAAATCTGTAGGATGAGAAAGATGTTGGGAAACGAAAAAGCCCTAGATAG 1560  
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Db 1561 AGAAAAAGATCTCTGAGTACAGTACTTATGGGGGGGGGGGAGGGGGGGATATCCACT 1620  
Qy 1621 GAGTCCAAAGTACTTGTGGGAGAGAAATCCACTGAGTACAAAGTACTTGTGGGGGAAGGAA 1680  
Db 1621 GAGTCCAAAGTACTTGTGGGAGAGAAATCCACTGAGTACAAAGTACTTGTGGGGGAAGGAA 1680  
Qy 1681 TGGCACAAGCAAAAGTTGAAGGGAAGAGAGATGGAGAGCCCTCAATGTTGGGGGTG 1740  
Db 1681 TGGCACAAGCAAAAGTTGAAGGGAAGAGAGATGGAGAGCCCTCAATGTTGGGGGTG 1740  
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Db 1741 TGAAGGTCACTCTCTCTTTTCCATGATGAGAGATTAAGAAAAATCAGTGTGTGAGTTG 1800  
Qy 1801 ATGTCCTTACACACCCCACTATGGCAGACTGTGGGAGACTTGGCATTTAGGGAAGGCGC 1860  
Db 1801 ATGTCCTTACACACCCCACTATGGCAGACTGTGGGAGACTTGGCATTTAGGGAAGGCGC 1860  
Qy 1861 GGCTTTTTCACACGAGAAACTTTATGCTCATCTCTGTGTGTACACTCCACCTTTGATGAG 1920  
Db 1861 GGCTTTTTCACACGAGAAACTTTATGCTCATCTCTGTGTGTACACTCCACCTTTGATGAG 1920  
Qy 1921 GTTAAGCTCAGGTTTCGTTCTACCGTTCTTGTCTACTGTGGGAACTTCACTAGTAGATTC 1980  
Db 1921 GTTAAGCTCAGGTTTCGTTCTACCGTTCTTGTCTACTGTGGGAACTTCACTAGTAGATTC 1980  
Qy 1981 CCAAGACAGGAGCAGCTCTTGTAAAGGAGGACCTGATTTCACTGTCTTAGAGAAC 2040  
Db 1981 CCAAGACAGGAGCAGCTCTTGTAAAGGAGGACCTGATTTCACTGTCTTAGAGAAC 2040  
Qy 2041 GAAATAGCTCAGAGAAATCTAGGTCAACGTAAGTCTAGGTCAACGCGGGCAAAATGACT 2100  
Db 2041 GAAATAGCTCAGAGAAATCTAGGTCAACGTAAGTCTAGGTCAACGCGGGCAAAATGACT 2100  
Qy 2101 GAAGCCTCTATTCCAGGTGAAACGCTGCTGATATATCTAGGTCAACGCGGGCAAAATGACT 2160  
Db 2101 GAAGCCTCTATTCCAGGTGAAACGCTGCTGATATATCTAGGTCAACGCGGGCAAAATGACT 2160  
Qy 2161 CACCGGATAAGATTTCTGTTAGTGTCTGCTTTTATTTTGCAGCACATCAGTGTGACGA 2220  
Db 2161 CACCGGATAAGATTTCTGTTAGTGTCTGCTTTTATTTTGCAGCACATCAGTGTGACGA 2220  
Qy 2221 CCAGAAATCCAGAGAAATCTCAGAGGCTGAAAGGAGACAGTGAAGGACTATTGGC 2280  
Db 2221 CCAGAAATCCAGAGAAATCTCAGAGGCTGAAAGGAGACAGTGAAGGACTATTGGC 2280  
Qy 2281 AAGCCAATACTAAGCCATTCAGTAGGAGACGTGGGGATTTCTTCTCTGCTTCCAGT 2340  
Db 2281 AAGCCAATACTAAGCCATTCAGTAGGAGACGTGGGGATTTCTTCTCTGCTTCCAGT 2340  
Qy 2341 CTCTTCTACTTTGTAAACATTTTCTTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2400  
Db 2341 CTCTTCTACTTTGTAAACATTTTCTTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 2400  
Qy 2401 CTGCACCTGATCTAGCTGGGTCTATAGATCTTTTCAATCTGTGTCTAAATTTGTAAGTCA 2460  
Db 2401 CTGCACCTGATCTAGCTGGGTCTATAGATCTTTTCAATCTGTGTCTAAATTTGTAAGTCA 2460  
Qy 2461 CAATTCTGGAGCTAGCAGAAAGCTTAGCTCAGCCAGCTCTCATGAGCACTTGTCTGGAGGA 2520  
Db 2461 CAATTCTGGAGCTAGCAGAAAGCTTAGCTCAGCCAGCTCTCATGAGCACTTGTCTGGAGGA 2520

Qy	2521	TGGCTTGTGACAGAGTCAATGCTTAAAGACAGCATCCCTGATTCCTCCAGCTCTCGCATTTGC	2581
Db	2521	TGGCTTGTGACAGAGTCAATGCTTAAAGACAGCATCCCTGATTCCTCCAGCTCTCGCATTTGC	2580
Qy	2581	CTAGTGGCCACGTGTAAATTACTTTTAGCTCGTAAATTAAGTATTTGGGAAAGCCAAATTTCCAC	2640
Db	2581	CTAGTGGCCACGTGTAAATTACTTTTAGCTCGTAAATTAAGTATTTGGGAAAGCCAAATTTCCAC	2640
Qy	2641	GACCTACATAATCCGAAGAAGCATGCAATGAAAACTAGAAAGCTGGGCACAAAATTTACTTA	2700
Db	2641	GACCTACATAATCCGAAGAAGCATGCAATGAAAACTAGAAAGCTGGGCACAAAATTTACTTA	2700
Qy	2701	GAGATGATTTTGGAGCTCATTAATACTGATGCTCTGAAATGTGATCAAAATCAACCCAGAA	2760
Db	2701	GAGATGATTTTGGAGCTCATTAATACTGATGCTCTGAAATGTGATCAAAATCAACCCAGAA	2760
Qy	2761	AACAACAAAAGAGCTGGATTTGCAAAATAGGACAAGTATTTAGAACTACTCGGTATTAACAG	2820
Db	2761	AACAACAAAAGAGCTGGATTTGCAAAATAGGACAAGTATTTAGAACTACTCGGTATTAACAG	2820
Qy	2821	CTGTCACTCTAAATTTAAATAATAGTGTCTTATTTAGCTGCCTATTTAAAGATTTAAACAACA	2880
Db	2821	CTGTCACTCTAAATTTAAATAATAGTGTCTTATTTAGCTGCCTATTTAAAGATTTAAACAACA	2880
Qy	2881	GTGGATTAACCTTCCAAATTTACTGGGCTCGTTCCTCAATAGAGTAAAAATATCAGTCACTAG	2940
Db	2881	GTGGATTAACCTTCCAAATTTACTGGGCTCGTTCCTCAATAGAGTAAAAATATCAGTCACTAG	2940
Qy	2941	TTAATTAATAGTGCATCAAGAGTATGAGTGGAAACCTTTCCCTTACTTTTAACTTCTTCA	3000
Db	2941	TTAATTAATAGTGCATCAAGAGTATGAGTGGAAACCTTTCCCTTACTTTTAACTTCTTCA	3000
Qy	3001	TCCTTAGTATTAATTTTCTTTCACACCTCGATCAAGCCACTAGTAAGCACCCTATCTG	3060
Db	3001	TCCTTAGTATTAATTTTCTTTCACACCTCGATCAAGCCACTAGTAAGCACCCTATCTG	3060
Qy	3061	CTGCGAGCTATTAATGACTTTACAGCAAAACAATTCGTGTGTGGCTCTTTGGGGAAG	3120
Db	3061	CTGCGAGCTATTAATGACTTTACAGCAAAACAATTCGTGTGTGGCTCTTTGGGGAAG	3120
Qy	3121	GGACAGGATAGCAGGAGGCTCAGGCTAGCAAGTCTGGACTCAACCTTAAAGCCAGAGGCA	3180
Db	3121	GGACAGGATAGCAGGAGGCTCAGGCTAGCAAGTCTGGACTCAACCTTAAAGCCAGAGGCA	3180
Qy	3181	TGGTTGATAGCAGAGAAAGTGAGGCTCTTCAAGTGGGTGCTGCTAAAGTAATCAGAAAC	3240
Db	3181	TGGTTGATAGCAGAGAAAGTGAGGCTCTTCAAGTGGGTGCTGCTAAAGTAATCAGAAAC	3240
Qy	3241	AGGAAGGCTCTGGTTGATGGAATTAATCAGTAAGATATCTACCTTATCTCTCTTCTAT	3300
Db	3241	AGGAAGGCTCTGGTTGATGGAATTAATCAGTAAGATATCTACCTTATCTCTCTTCTAT	3300
Qy	3301	AGAAGCTTAAACCGTCTCTCTTCTTGTGTAGGCTGATAAACACGCTTGTTTTCTTTTG	3360
Db	3301	AGAAGCTTAAACCGTCTCTCTTCTTGTGTAGGCTGATAAACACGCTTGTTTTCTTTTG	3360
Qy	3361	AGTGTTCATGGCTTTGCAGATTTTTCAGTGTCTGCCAGTTCCTTGTAGAGGGTTTGTTC	3420
Db	3361	AGTGTTCATGGCTTTGCAGATTTTTCAGTGTCTGCCAGTTCCTTGTAGAGGGTTTGTTC	3420
Qy	3421	CTTGACACCTGGGCTTGGATGTTTAGCATGCAAGGACACACCTTCGTAAATGCTGTGTA	3480
Db	3421	CTTGACACCTGGGCTTGGATGTTTAGCATGCAAGGACACACCTTCGTAAATGCTGTGTA	3480
Qy	3481	AAAGGTTATTATTCATTACTTTTGTCTTTGGAAAGGTGAAGTGTGTGAGAAAGAACCTC	3540
Db	3481	AAAGGTTATTATTCATTACTTTTGTCTTTGGAAAGGTGAAGTGTGTGAGAAAGAACCTC	3540
Qy	3541	ACAGGAGATGTATTCCTCTGTAGGAAACCTTTTTTTTCCCTTAAAGCCCTATAATCCACT	3600
Db	3541	ACAGGAGATGTATTCCTCTGTAGGAAACCTTTTTTTTCCCTTAAAGCCCTATAATCCACT	3600
Qy	3601	TTCACTCAACTTTGACTTTTATACCATGCTGTGCATGAGAAAGAGTGTTTAGGCCCGCTCT	3660

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3661	CGT	GGCT	CT	CGG	AAAG	CAC	CAAT	TAG	GGG	AAAG	AAT	TGTT	TAT	CCG	CAG	AAAT	CT	GACT	GGC	3720							
3661	CGT	GGCT	CT	CGG	AAAG	CAC	CAAT	TAG	GGG	AAAG	AAT	TGTT	TAT	CCG	CAG	AAAT	CT	GACT	GGC	3720							
3721	AGG	AAA	CT	GGT	CAG	AGCT	CCCC	AAAG	ACC	ACT	CAC	AGT	GT	TAA	GT	TAG	GAA	CAGT	CGAG	3780							
3721	AGG	AAA	CT	GGT	CAG	AGCT	CCCC	AAAG	ACC	ACT	CAC	AGT	GT	TAA	GT	TAG	GAA	CAGT	CGAG	3780							
3781	GGT	GGT	TC	ATA	TAT	TAG	AAT	TGGA	AC	GAG	GGG	AGG	AG	AT	TA	AGCT	AC	AAAGT	TT	CA	TAG	3840					
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3841	GGT	CT	CA	TAGT	CT	TT	TAA	GAT	ACAA	AT	TAG	CT	TGGT	TT	CA	TAA	CAAG	AG	AGT	CT	TGGG	3900					
3901	AAG	G	CAG	CA	AGCA	TT	GAG	AGG	AGAT	TGG	AAAGG	AAAA	CAAT	GT	TAG	AGG	AT	TTG	AAAA	3960							
3901	AAG	G	CAG	CA	AGCA	TT	GAG	AGG	AGAT	TGG	AAAGG	AAAA	CAAT	GT	TAG	AGG	AT	TTG	AAAA	3960							
3961	GCT	ACAA	T	CCCT	CC	CAG	AGG	AGT	TTTT	TT	CT	TGG	AG	GAAT	CT	TAG	AA	CAAG	GGT	GGT	G	4020					
3961	GCT	ACAA	T	CCCT	CC	CAG	AGG	AGT	TTTT	TT	CT	TGG	AG	GAAT	CT	TAG	AA	CAAG	GGT	GGT	G	4020					
4021	AGG	T	CGAT	TC	G	CAG	AGG	AGCT	TT	GCT	TT	GC	CAAT	TT	GAAT	CT	T	GC	T	CC	AT	TGA	4080				
4021	AGG	T	CGAT	TC	G	CAG	AGG	AGCT	TT	GCT	TT	GC	CAAT	TT	GAAT	CT	T	GC	T	CC	AT	TGA	4080				
4081	GGT	T	GAG	AG	CGCT	CAC	CCCT	TTTT	TAC	CCCT	GGAT	GAG	AGG	AAAA	CAAG	GGG	T	GT	TTT	TG	AC	4140					
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4141	TCCT	AC	CT	G	AGT	TTTT	T	ACT	AGT	TTTA	CG	CAAT	TG	GAAC	CAG	AC	CT	CG	G	AC	CT	CT	CT	TG	AC	4200	
4201	P	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	4260
4201	P	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	4260
4261	GCA	CAGG	CAG	CT	GGG	CA	TGGT	GGCC	AT	GC	CTTT	TAAT	CC	CAG	CA	TTT	GGG	AGG	CAG	AGGC	4320						
4261	GCA	CAGG	CAG	CT	GGG	CA	TGGT	GGCC	AT	GC	CTTT	TAAT	CC	CAG	CA	TTT	GGG	AGG	CAG	AGGC	4320						
4321	AGG	T	GACT	TTT	CT	AAAT	TC	AA	GGCC	AG	CT	TGGT	CT	CA	AA	GT	AG	TT	C	CAG	CA	CG	CAGG	4380			
4321	AGG	T	GACT	TTT	CT	AAAT	TC	AA	GGCC	AG	CT	TGGT	CT	CA	AA	GT	AG	TT	C	CAG	CA	CG	CAGG	4380			
4381	GCT	AT	AC	G	AG	AA	AC	CT	GT	CT	CGG	AA	AAAA	AAAA	AAAA	AG	AA	AAAA	AAAA	AAAA	AG	AA	AG	4440			
4381	GCT	AT	AC	G	AG	AA	AC	CT	GT	CT	CGG	AA	AAAA	AAAA	AAAA	AG	AA	AAAA	AAAA	AAAA	AG	AA	AG	4440			
4441	ACA	G	CAG	CAG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	4500						
4441	AGA	G	CAG	CAG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	4500						
4501	AGG	AG	GAG	GAG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	4560						
4501	AGG	AG	GAG	GAG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	4560						
4561	PAG	ACA	GAG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	AGG	4620						
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4621	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	AGAAA	4680						
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4681	GGT	CTTT	GAG	ACA	AGG	CTTT	TG	AA	TTG	AG	CGCT	TC	AT	CA	AT	AGT	TG	AT	CA	TG	GGT	CAG	TGG	4740			

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Qy 4741 AGGCTACCTGCTCAGCGCGAGCCCTGCTGCGTTAGCACTTAACTCTCCAGGTCTCAGTA 4800

Db 4741 AGGCTACCTGCTCAGCGCGAGCCCTGCTGCGTTAGCACTTAACTCTCCAGGTCTCAGTA 4800

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Qy 4861 AAAATTTAAATTTACAAAAGGCGAGTGTATATTTGTGGGATACAGTGTGATTAATGATGATG 4920

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Db 5101 TTTTAGAAGGCTGATCTTGGTTTGGTGTCTCAGCAAGCAAAATGTCACAGCTCTTTC 5160

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Db 5281 TCTGAGAAATGCTTGCCTGCTGAGGAGAAAGCTAGAAAACGAAGACTGCTCCTTCCT 5340

Qy 5341 GCCTTCTAAAAGAACATAAGATCCCTGGAATGACTTTTTTACTTAAAGGAAAGTGAGAA 5400

Db 5341 GCCTTCTAAAAGAACATAAGATCCCTGGAATGACTTTTTTACTTAAAGGAAAGTGAGAA 5400

Qy 5401 GCTAACGTCCTCCATCATTAGAAGATTTCAATGAAACCTGGCTCAGTTGAAAGAGAAA 5460

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Db 5881 TTACCCAATCATTCTCATGACTTCATGCTTGACTCATATATTATCTGGTAAAGTTTG 5935

RESULT 2

US-09-419-568F-29

Sequence 29, Application US/09419568F

Patent No. 6331613

GENERAL INFORMATION:

APPLICANT: Dumoutier, Laure

APPLICANT: Renauld, Jean-Christophe

TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

FILE REFERENCE: LUD 5543.2

CURRENT APPLICATION NUMBER: US/09/419,568F

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PRIOR FILING DATE: 1999-07-16

PRIOR APPLICATION NUMBER: US09/178,973

PRIOR FILING DATE: 1998-10-26

NUMBER OF SEQ ID NOS: 29

SEQ ID NO 29

LENGTH: 5935

TYPE: DNA

ORGANISM: Mus musculus

FEATURE:

US-09-419-568F-29

Query Match 100.0%; Score 5935; DB 3; Length 5935;

Best Local Similarity 100.0%; Pred. No. 0;

Matches 5935; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1 GAATTCGAAGTCCACATCGAATCAATCCGAATACCTTTGTAATTTCTCTTCTTCAAAATATCC 60

Qy 61 ATCTATATAGTATAGTTATTTAGGATCATTTAAAAATATATGTTTGGAGACTTATGTTT 120

Db 61 ATCTATATAGTATAGTTATTTAGGATCATTTAAAAATATATGTTTGGAGACTTATGTTT 120

Qy 121 GCACAAGTAAATGTCAGAGAGAAATTAGCAAAATGATATATTATTTTAAAAAAT 180

Db 121 GCACAAGTAAATGTCAGAGAGAAATTAGCAAAATGATATATTATTTTAAAAAAT 180

Qy 181 CTATGCTTAAATGCTATTAGATTGTTCACTACTGACATTTCCAAACTTAACTTGACCT 240

Db 181 CTATGCTTAAATGCTATTAGATTGTTCACTACTGACATTTCCAAACTTAACTTGACCT 240

Qy 241 TGGCTATGATTTCAACCTTTGTTATTTGCATCTACCAATACTGTGTCTCACTTACCATGC 300

Db 241 TGGCTATGATTTCAACCTTTGTTATTTGCACTACCAATACTGTGTCTCACTTACCATGC 300

Qy 301 TATCCGACGAGCATGTTCCCTGATGTTTTTGGCTTTTGGCTCTCTCGCTAACAGGCTCTC 360

Db 301 TATCCGACGAGCATGTTCCCTGATGTTTTTGGCTTTTGGCTCTCTCGCTAACAGGCTCTC 360

Qy 361 CTCTCAGTTATCAACTTTTGACACTTTGCGATCGGTGATGGCTGCTCGAGAAATCTA 420

Db 361 CTCTCAGTTATCAACTTTTGACACTTTGCGATCGGTGATGGCTGCTCGAGAAATCTA 420

Qy 421 TGAGTTTTTCCCTTATGCGGACTTTTGGCCGCGACCTGCTCTCTCTCATTCGCCCTGTGG 480

Db 421 TGAGTTTTTCCCTTATGCGGACTTTTGGCCGCGACCTGCTCTCTCTCATTCGCCCTGTGG 480

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Db 541 AGCAGCGTATACGTCAACCGCACCTTTATGTGGCCAAAGGAGTACAGCTCATCTCT 600  
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Db 1801 ATGCTTTCAGACACCCCAACTATGGCAGACTGTGGAGACCTGGCAATTTAGGAAGCGCG 1860  
Qy 1861 GGCCTTTTACAGAGAAACTTTATGCTCATCTCTGTGTCTACACTCCACCTTTTATGATG 1920  
Db 1861 GGCCTTTTACAGAGAAACTTTATGCTCATCTCTGTGTCTACACTCCACCTTTTATGATG 1920  
Qy 1921 GTTAAAGCTCAGGTTTCTGTTTCTACCGTCTTGTCTACTGGTGGAAAATTTCACTAGGATTC 1980  
Db 1921 GTTAAAGCTCAGGTTTCTGTTTCTACCGTCTTGTCTACTGGTGGAAAATTTCACTAGGATTC 1980  
Qy 1981 CCAAGACGAGGACAGCTCTTCTGTAAAGGAGGACCTGGATTTCACTGTCTTAGAGAAC 2040  
Db 1981 CCAAGACGAGGACAGCTCTTCTGTAAAGGAGGACCTGGATTTCACTGTCTTAGAGAAC 2040  
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Qy 2161 CACCGGATAAGATTTCTGTTAGTGAGTCTGCTTTTATTTTGCAGCACATCAGTGGTACGA 2220  
Db 2161 CACCGGATAAGATTTCTGTTAGTGAGTCTGCTTTTATTTTGCAGCACATCAGTGGTACGA 2220  
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Db 2521 TGGCTTGTGACAGAGTCAATGCTTAGAAGACAGCATCCCTGATTTCCAGCTCTGCACTTGC 2580  
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Db 2581 CTAGTGGCCACGTTGTAATTTACTTTTAGCTGATTAAGTATTTGGGAAGGCAATTTCCACC 2640  
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Qy	2701	GAGATGATTTTGGAGCTCATTAACACTGATGCTCTGAAATGTGATCAAAATCAACCAGAA	2760
Db	2701		2760
Qy	2761	AACAAACAAAGAGCTGGATTTGC AAAATAGCAAAAGTATTTAGAAATCACTGGTATTAACAG	2820
Db	2761		2820
Qy	2821	CTGTCATCTTAATTAATAATATAGTGTCTATTTAGCTGCCTATTTAAGATTAACACACAAGA	2880
Db	2821		2880
Qy	2881	GTGGATAACTCCCAATTTACTCGGCCTGGTTTCAATAGAGTAAATAATATCAGTCAATAGA	2940
Db	2881		2940
Qy	2941	TTAATTAATAGTGTGATGAAGTATGAGTGTGAAAACCCCTTCTCTTACTTTTACCTTCATT	3000
Db	2941		3000
Qy	3001	TCCTTAGTATTAATTTTCTTCTTCAACCCCTGATCAAGCCACTAGTAAGCACCTATCTG	3060
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Qy	3061	CTCGAGACTATATATGACTTTTACAGCAAAACAATGCTGTGTGGCTCTTTGGGGAAG	3120
Db	3061		3120
Qy	3121	GGAAACAGGATAGCAGGAGGCTCAGGCTAGCAAGTCTGAGCTCAACCTAAAGCCAGAGGCA	3180
Db	3121		3180
Qy	3181	TGGTTGATAGCAGAGAAAGTGAAGCTCTTCAAAAGTGGGTGTGCTTAAGTAATCAGAAAC	3240
Db	3181		3240
Qy	3241	AGGAAGGCTCTGGTTGATGGAATTAATCAGTAAGATATCTACCCCTATCTCCTCTCTTAT	3300
Db	3241		3300
Qy	3301	AGAAAGCTAAACCGTCTCTCTTCTTGTGTGTAGGCTGATAAACACGCTGTGTTCTTTTG	3360
Db	3301		3360
Qy	3361	AGTGTTCATGCTTTGAGATTTTCAGTGTCTGCCAGTCTCTGTAGAGGGTTGTGTAC	3420
Db	3361		3420
Qy	3421	CTTGACACTGGCTTGGATGTTAGCATGCCAAAGGCCACACACTTCTGAATGCCCTGTGTA	3480
Db	3421		3480
Qy	3481	AAAGGTTATTATTCATTTTACTTTGTCTTTGGAAAAGGTGAAGTGTGTGGAAGAACTC	3540
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Qy	3541	ACAGAGATGATTTCTGTGAGAAAACCTTTTTTTTCCCTTAAAGCCCTATATCCACT	3600
Db	3541		3600
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Qy	3661	CGTGCTCTGGGAAAAGCACAATAGGGGAAGAAATGTTATGCCGAGAAATCTGACTGGC	3720
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Qy	3721	AGGGAAACTGGGTCTAGAGCTCCCAAGCACTACAGGTGTTAAGTAGGAACAGTCCAG	3780
Db	3721		3780
Qy	3781	GGTGGGTTCAATATAAGTAAGTAAGGAGGGAAGATAAGCTACAAAGTTTCATAG	3840
Db	3781		3840

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Qy	3841	GGTCCTTAAGTCTTTAAGATACAAAATAGCTGGTTGGCTTCAATAAAGGAAAGTCTGGG	3900
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Qy	4021	AGGTGGATCCAGAGAGCACTTGTTCGCAATTTGAAATCTGGGTTTTTGTCTCTCCATTGA	4080
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Qy	4141	TCCTACCTGGAGTTTACTAGTTTACGCAATGGAACACACACTCGGGACCTCTCTTTGAC	4200
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Qy	4201	AGAAAAAAGGAAACCTGTGTTTCTCTGTTGTTCTCTGTTGTTTGTAAAGAA	4260
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Qy	4261	GCACAGCAGCTGGCATGTGGCCCATGCTTTAATCCAGCAATTTGGGAGGCGAGGC	4320
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Qy	4321	AGGTGACTTTCTAAATTTCAAGGCCAGCTGGTCTACAAAAGTGAGTTCCAGGACAGCCAGG	4380
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Qy	4441	AGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAG	4500
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Qy	4501	AGGAGGAG	4560
Db	4501		4560
Qy	4561	AG	4620
Db	4561		4620
Qy	4621	AGAAAG	4680
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Qy	4801	TCACCTTCTGCTCTAGCAGTTAGGAGTTAGCAAAACCTTTTTTCCAAACCCCACT	4860
Db	4801		4860
Qy	4861	AAAAATTTAATTAACAAAGGAGGAGTAAATTTGTGGGATACAGTGTGATTAATTTGAT	4920
Db	4861		4920



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QY 2701 GAGATGATTTTGTAGCTCATTTAAACTGATGCTCTGAAATGTGATCAAAATCAACCCAGAA 2760  
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; Sequence 8, Application US/09178973B
; Patent No. 6274710
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renaud, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: (TIFs)
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543
; CURRENT APPLICATION NUMBER: US/09/178.973B
; CURRENT FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 17
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; SEQ ID NO 8
; LENGTH: 7445
; TYPE: DNA
; ORGANISM: Mus musculus
US-09-178-973B-8
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Query Match 71.5%; Score 4245.2; DB 3; Length 7445;
Best Local Similarity 88.3%; Pred. No. 0;
Matches 5039; Conservative 0; Mismatches 178; Indels 487; Gaps 20;
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; Sequence 8, Application US/09419568F  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/419,568F  
; CURRENT FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-419-568F-8

Query Match 71.5%; Score 4245.2; DB 3; Length 7445;  
Best Local Similarity 88.3%; Pred. No. 0;  
Matches 5039; Conservative 0; Mismatches 178; Indels 487; Gaps 20;

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Db 7245 GTG 7248

RESULT 6  
US-09-354-243B-8  
; Sequence 8, Application US/09354243B  
; Patent No. 6359117  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fa  
; TITLE OF INVENTION: (TIFs)  
; FILE REFERENCE: LUD 5543.1  
; CURRENT APPLICATION NUMBER: US/09/354,243B  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-354-243B-8

Query Match 71.5%; Score 4245.2; DB 3; Length 7445;  
Best Local Similarity 88.3%; Pred. No. 0;  
Matches 5039; Conservative 0; Mismatches 178; Indels 487; Gaps 20;

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Db 3347 GCATGGAGATCCACTGAGTACAAAGTACTTGTGGGGGGAGGAAATGGCAGAGCAAGAAAGT 3406

QY 1698 TGAAGGAAAGAGAGAAATGGAGAGGCTCAATGTTGGGGGTGTGAAGGTCACTCTT 1757  
Db 3407 TGAAGGAAAGAGAGAAATGGAGAGGCTCAATGTTGGGGGTGTGAAGGTCACTCTT 3462

QY 1758 TTCCATGTGATGAGAGTTAAGAAATCACTGTGTGATTTGATGCTTCTCAGACACCC 1817  
Db 3463 TTCCATGTGATGAGAGTTAAGAAATCACTGTGTGATTTGATGCTTCTCAGACACCC 3522

QY 1818 AA-----CTATGGCAGCTGTGGGAGACCTGGCAATTTAGGAA 1855  
Db 3523 CAACATATGAACATATCCACGAGGAGGGGAGACTGTGGGAGACCTGGCAATTTAGGAA 3582

QY 1856 GGGCGGCTTTTTCAGAGAAATTTTATGCTCTCTTGTGCTACACTCCACCTT 1915  
Db 3583 GGGCGGCTTTTTCAGAGAAATTTTATGCTCTCTTGTGCTACACTCCACCTT 3642

QY	1916	ATGAGGTTAAGCTCAGGTTTCGTTTCTACCGTTCTTGTCTACTGCTGGAACCTTCAGTAGG	1975
Db	3643	ATGAGGTTAAGCTCAGGTTTCGTTTCTACCGTTCTTGTCTACTGCTGGAACCTTCAGTAGG	3702
QY	1976	ATTCCCAAAAGACGAGGACAGCTCTCTGTGAAGGAGGACCTGGATTTCAGTGTCTCAG	2035
Db	3703	ATTCCCAAAAGACGAGGACAGCTCTCTGTGAAGGAGGACCTGGATTTCAGTGTCTCAG	3762
QY	2036	AGAACGAAATAGCTCAGAGAATCTAGGTCAACGTGAATCTAGGTCAACGCGGCAAAAA	2095
Db	3763	AGAACGAAATAGCTCAGAGAATCTAGGTCAACGTGAATCTAGGTCAACGCGGCAAAAA	3822
QY	2096	TGACTGAACCGCTCTATTCCAGGTGAACGGTCAAGTGCCTCAGATATACGTAGGTATTGG	2155
Db	3823	TGACTGAACCGCTCTATTCCAGGTGAACGGTCAAGTGCCTCAGATATACGTAGGTATTGG	3882
QY	2156	GCTCCCAACCGGATAGGATCTGTGTAGTGTCTGTCTTTTATTTTGCAGCACATCAGTGGT	2215
Db	3883	GCTCCCAACCGGATAGGATCTGTGTAGTGTCTGTCTTTTATTTTGCAGCACATCAGGCGT	3942
QY	2216	GACGACAGAACATCCAGAGAATGTCCAGAGGCTGAAGGAGACAGTGAAAAAGGTACTA	2275
Db	3943	GACGACAGAACATCCAGAGAATGTCCAGAGGCTGAAGGAGACAGTGAAAAAGGTACTA	4002
QY	2276	TTGGCAAGCCACAATACTAAGCCATTTCAGTAGGAGACGTGGGGATTTCTTCTCTGCTTC	2335
Db	4003	TTGGCAAGCCACAATACTAAGCCATTTCAGTAGGAGACGTGGGGATTTCTTCTCTGCTTC	4062
QY	2336	CCAGTCTCTCTACTTTTGAACATTTCTTTGTAGTGTCTACTGTCTGCTGGTCCATTACTCA	2395
Db	4063	CCAGTCTCTCTACTTTTGAACATTTCTTTGTAGTGTCTACTGTCTGCTGGTCCATTACTCG	4122
QY	2396	CTTAGCTGCACCTGCATCTAGCTGGGTCTATAGATCTTTCAATCTGTGTCTAAATTTGTA	2455
Db	4123	CTTAGCTGCACCTGCATCTAGCTGGGTCTATAGATCTTTCAATCTGTGTCTAAATTTGTA	4182
QY	2456	AGTCACAATTCGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTCATGAGACATTGCTCG	2515
Db	4183	AGTCACAATTCGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTCATGAGACATTGCTCG	4242
QY	2516	GAGGATGGCTGTGACAGAGTCAATGCTAGAGACAGCATCCCTGATTTCCAGCTCTGCA	2575
Db	4243	GAGGATGGCTGTGACAGAGTCAATGCTAGAGACAGCATCCCTGATTTCCAGCTCTGCA	4302
QY	2576	CTTGCTCTAGTGGCCACCGTAAATTTACTTTTACCTGTGATTAAGTATTTGGGAAAGCCAACTC	2635
Db	4303	CTTGCTCTAGTGGCCACCGTAAATTTACTTTTACCTGTGATTAAGTATTTGGGAAAGCCAGTTC	4362
QY	2636	CCACCGACCTACATAATCCGAAGAGCATGCAATTTGAAAACTAGAAAGCTGGGCACAAACT	2695
Db	4363	CCACCGACCTACATAATCTGAAGAACCAATGCAATTTGAAAACTAGAAAGCTGGGCACAAACT	4422
QY	2696	TACTAGAGATGATTTTGAAGTCAATTAACCTAGTGTCTGAAATGTGATCAATCAACCC	2755
Db	4423	TACTAGAGATGATTTTGAAGTCAATTAACCGAGTGTCTGAAATGTGCGCAAAATCAACCC	4482
QY	2756	AGAATAACAAAGAGCTGGATTTCGAAATAGGACAGTATTTAGAATCACTGGTATT	2815
Db	4483	AGAATAACAAAGAGCTGGATTTCGAAATAGGACAGTATTTAGAATCACTGGTATT	4542
QY	2816	AACAGCTGTCTATTAATTAATAATATAGTGTCTATTTAGTCTGCTATTTAAGATTAAACA	2875
Db	4543	AATAGCTATCATCTTAATTAATAATATAGGCTTATATA --- TATATTTAAGATTAAACA	4598
QY	2876	CAAGAGTGGATTAATCTCCCAATTTACTGGGCTGGTTTCAATAGAGTAAAAAATATCAGTC	2935
Db	4599	CAAGAGTGGATTAATCTCCCAATTTACTTGGCTGGTTTCAAAAGAGTAAAAAATATCAGTC	4658
QY	2936	ATAGATTAAATTAATAGTGTCAATGAAGTATGAGTTGGAAACCTTTTCTTACTTTTACCT	2995
Db	4659	ATAGATTAAATTAATAGTGTCAATGAAGTATGAGTTGGAAACCTTTTCTTACTTTTACCT	4718
QY	2996	TCATTTCTTAGTTATTTATTTTCTTCTCACACCTTGATCAAGCCACTAGTAGACACCT	3055

Db	4719	TCATTTCTTAGT --- TTTTCTTCTCACACCTTGATCAAGCCACTAGTAGACACCT	4773
QY	3056	ATCTGCTGCAGCTATTATATGACTTTTACAGCAAAACAATTTGCTGTGTGGCTCTTTGG	3115
Db	4774	ATCTGCTGCAGCTATTATATGACTTTTACAGCAAAACAATTTGCTGTGTGGCTCTTTGG	4833
QY	3116	GGAAGGGAACAGGATAGCAGGAGGCTCAGCTAGCAAGTCTGGAATCAACTAAAGCCAG	3175
Db	4834	GGAAGGGAACAGGATAGCAGGAGGCTCAGCTAGCAAGTCT - GACTTGGCCCTAAAGCCAG	4892
QY	3176	AGGCATGGTTGATAGCAGAGAAAGTGGGCTCTTCAAAAGTGGGTGGCTTTAAGTAATCA	3235
Db	4893	AGGCATGGTTGATAGCAGAGAAAGTGGGCTCTTCCAAAGTGGGTGGCTTTAAGTAATCA	4952
QY	3236	GAAACAGGAAGGCTCTGGTTGATGGAAATTAATCAAGTAAGATATCTACCTTATCTCTCT	3295
Db	4953	GAAACAGGAAGGCTCCGGTTGATGGAAATTAATCAAGTAAGATATCTACCTTATCTCTCT	5009
QY	3296	TCATAGAAAGCTAAACCGTCTCTCTTTGTGTAGGCTGATAAACAACGCTTTGTTTC	3355
Db	5010	TCATAGAAAGCTAAACCGTCTCTCTTTTCTTGTGTAGGCTGATAAACAACGCTTTGTTTC	5069
QY	3356	TTTGTAGTGTTCATGGCTTTGCAAGATTTTTCAGTGTCTCTGCCAGTCTTGTGTAGAGGTTT	3415
Db	5070	TTTGTAGTGTTCATGGCTTTTGTAGATTTTGTAGTGTCTCTGCCAGTCTTGTGTAGAGGTTT	5129
QY	3416	GTGTACCTTTGACACCTGGGCTTGGATGTTTACATGTTAGCATGCCAAGGACACACTTCTGAATGCCT	3475
Db	5130	GTGTACCTTTGACACCTGGGCTTGGATGTTTACATGTTAGCATGCCAAGGACACACTTCTGAATGCCT	5189
QY	3476	GTGTAAAAGGTTTATTTCAATTTTGTCTTTTGGAAAGGTGAAGTGTGTGAGAAAG	3535
Db	5190	GTGTAAAAGGTTTATTTCAATTTTGTCTTTTGGAAAGGTGAAGTGTGTGAGAAAG	5249
QY	3536	AACCTCAGAGAGATGATTTCTCTGTAGGAAAC - TTTTCTTCTCCCTTAAAGCCTATAA	3594
Db	5250	AACCTCAGAGAGATGATTTCTCTGTAGGAAACCTTTTCTTCTTAAAGCCTATAA	5309
QY	3595	TCCACTTTTCACTCACTTTTACATGCTGTACATGAAAGAGTGTGTAGGCC	3654
Db	5310	TCCACTTTTCACTCACTTTTACATGCTGTACATGAAAGAGTGTGTAGGCC	5369
QY	3655	CGCTCTCTGGCTCTGGGAAAGCACCAATAGGGGAAAGAAATTTATSCCGAGAAATCTG	3714
Db	5370	CGCTCTCTGGCTCTGGGAAAGCACCAATAGGGGAAAGAAATTTATGCTGAGAAATCTG	5429
QY	3715	ACTGGCAGGGAAACTGGGTCAGAGCTCCCAAAAGACCACTACAGGTTTAAAGTAGGAACA	3774
Db	5430	ACCGCAGGGAAACTGGTCAGAGCTCCCGGAAAGACCACTACAGGTTTAAAGTAGGAACA	5489
QY	3775	GTGAGGGTGGGTTTATTAATAGAAATGGAACAGAGGGGGAAGATAAGCTACAAAGTT	3834
Db	5490	GTGAGGGTGGGTTTATTAATAGAAATGGAACAGAGGGGGAAGATAAGCTACAAAGTT	5549
QY	3835	TCATAGGCTCCTTAAGTCTTTAAGATACAAATAGCTTTGGGCTTCATTAACAAAGGAAG	3894
Db	5550	TCATAGGCTCCTTAAGTCTTTAAGATACAAATAGCTTTGGGCTTCATTAACAAAGGAAG	5609
QY	3895	TCGGAAGGACGACGATTTGAGAGGAGATGGAAGGGAAGGAAAC - AATGTAGAGGA	3952
Db	5610	TCGGAAGGACGACGATTTGAGAGGAGATGGAAGGGAAGGAAACAGATGTAGAGGA	5666
QY	3953	TTTGAAGAGCTACAAATCTCCACGAGAGGATTTTCTTGGAGGAACTAGAACAAAGGTT	4012
Db	5667	CTTGAAGAGCTACAAATCTCTTACAGAGATTTTCTTGGAGGAACTAGAACAAAGGTT	5722
QY	4013	GGTGGATTAGGTCGACAGAGGACTCTCTTTTGGCATTTGAATCTGGGTTTTTGTCTC	4072
Db	5723	AGTGGATTAGT - GATTGACAGGGGACTTCTTTTGGCATTTGAATCTGGGTTTTTGTCTC	5781
QY	4073	TCATTTGAGGTTGAGAGCGTCACTTTTCTTACCTGGATAGGAGGAGGAAGAGGTT	4132

Db 5782 TCCATTGAGGTGAAAGCGTCAACC-TTTTTCACCTCGAATGGAGGAGAAAGAGGGGT 5840  
Qy 4133 GTTTTGACTCCTACTCGAGTTTACTAGTTTACGCAATGGAACAGACACTCGGGACCTC 4192  
Db 5841 GTTATGACTCCTACTCGAGTTTACTAGTTTACGCAATGGAACAGACACTCGGGACCTC 5900  
Qy 4193 CTCTTTGACAAAGAAAAAAGAAAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAG 4252  
Db 5901 CTCTTGAC-----AAAAAATGGAACCTGTGTTGTTGTTGTTGTTGTTGTTGTTG 5950  
Qy 4253 TTAAGAAAGCACAGCGAGCTGGGCACTGGGCACTGGGCCCATGCCCTTTAATCCAGCATTTGGGAG 4312  
Db 5951 TTAAGAAAGCACA----- 5963  
Qy 4313 GCAGAGGAGGTGACTTTCTAAATTTCAAGGCCACCTGGTCTACAAAGTGAGTTCCAGGA 4372  
Db 5964 ----- 5963  
Qy 4373 CAGCCAGGCTATACAGAGAAACCTGTCTCGGGAAGAAAAAGAAAGAAAGAAAG 4432  
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Qy 4433 AAAAGAGAGAGAGGAGAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 4492  
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Qy 4493 GAA 4552  
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Qy 4553 GAGAAGAA 4612  
Db 5964 ----- 5963  
Qy 4613 AAGAGAAAGAAAAAGAAAAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCA 4672  
Db 5964 -----GGCAAGCCCGACCAATGGGT 5985  
Qy 4673 CGTATGGGTCTTTGAGACAAGGCTTTTGAATTTGAGGGCTCATCAATAGTTGATCATGG 4732  
Db 5986 TGAATGTGGGTCTTTGAGTCAAGGCTTTTGAAGTTGAGCACTCATCAATAGTTGATCATGG 6045  
Qy 4733 TCAGGTGAGGGCTACCTGTGAGCGGAGCCCTGCTGCTTGTAGCACTTAAACATCTCCAGG 4792  
Db 6046 TCAGGTGAGGGCTACCTGTGAGCGGAGCCCTGCTGCTTGTAGCACTTAAACATCTCCAGG 6105  
Qy 4793 TCTCAGTATCACTTCTGCTGCTTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTCCAA 4852  
Db 6106 TCTCAGTATCACTTCTGCTGCTTTAGCACAGTTAGGAGTTGAGCAAAACCTTTTTTCCAA 6165  
Qy 4853 CCCCCACTAAATTTAATTTACAAAGCGCAGTGTAAATTTGGGATACAGTGTGATTAAT 4912  
Db 6166 CCCCCACTAAATTTAATTTGACAAAGACTGTGTAATTTGGGATACAGTGTGATTAAT 6225  
Qy 4913 GATCTATGTGTCATTGTGCAAGGTTCAATAAGGTAGATCAATAGGCCCATCAACAGCTT 4972  
Db 6226 GATCTATGTGTCATTGTGCAAGGTTCAATAAGGTAGATCAATAGGCCCATCAACAGCTT 6285  
Qy 4973 TATGGGTGTAAGTCAAGTAAATATAGGTAGATGCTGT-GTGTCTTTAGGTCTCAGAAGG 5031  
Db 6286 TATGGGTGTAAGTCAAGTAAATATAGGTAGATGCTGTGTGTGCTTTAGGTCTCAGAAGG 6345  
Qy 5032 CATGATTTTAAGGTCTGGGCAAT 5091  
Db 6346 CATGATTTTAAGGTCTGGGCAAT 6405  
Qy 5092 TTATCAATCTTTTAGAGAGGCTGATCTGTTTGGTGTCTCAGCAGCAAGCAAGTGTACCC 5151  
Db 6406 TTATTAATCTTTTAGAGAGGCTGATCTGTTTGGTGTCTCAGCAGCAAGCAAGTGTACCC 6465  
Qy 5152 AGCTCTTTCTAACTAGTACCACTTTTAGAAAAATGCTACCGGTCTCAAAATTTGGTTGTATT 5211  
Db 6466 AGCTCTTTCTAACTAGTACCACTTTTAGAAAAATGCTACCTGTGCTCAAAATTTGGTTGTATT 6525

Qy 5212 CTTATTTTTCATAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCT 5271  
Db 6526 CTTATTTTTCATAGCTTGGAGAGAGTGGAGAGATCAAGCCGATTTGGGAACTGGACCTGCT 6585  
Qy 5272 GTTTATGTCTCTGAGAAATGCTTTCGCTCTGAGCGAGAGAAAGCTTAGAAAAAGAAAGAACTG 5331  
Db 6586 GTTTATGTCTCTGAGAAATGCTTTCGCTCTGAGCGAGAGAAAGCTTAGAAAAAGAAAGAACTG 6645  
Qy 5332 CT 5391  
Db 6646 CT 6705  
Qy 5392 AAGTGAGAACTAAGCTCCACCATCATTTAGAAAGATTTCAATGAAAACTGGCTCAGTTGA 5451  
Db 6706 AAGTGAGAACTAAGCTCCACCATCATTTAGAAAGATTTCAATGAAAACTGGCTCAGTTGA 6765  
Qy 5452 AAGAGAAAAATAGTGTCAAGTTGTCCATGAGACCAAGAGTGTAGATAACCAAGAAAT 5511  
Db 6766 AAAAGAAAAATAGTGTCAAGTTGTCCATGAGACCAAGAGTGTAGATAACCAAGAAAT 6825  
Qy 5512 TCATTGCAATATTTTATTTGTCATTGATTAATGCAAGAAAAAGTATGTACTTTTAAAAAA 5571  
Db 6826 TCATTGCAATATTTTATTTGTCATTGATTAATGCAAGAAAAAGTATGTACTTTTAAAAAA 6885  
Qy 5572 TTGTTTGAAGAGGTTTACCTCTCATTCCTCTAGAGAAAAAGCCCTATGTAACTTTCATTTT 5631  
Db 6886 TTGTTTGAAGAGGTTTACCTCTCATTCCTCTTGTAGAAAAAGCTTATGTAACTTTCATTTT 6945  
Qy 5632 CATAAACCAATCTTTATATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTC 5691  
Db 6946 CATATCCAATATTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTC 7005  
Qy 5692 AGTTTATTAATATGGAATTTTATATAGAAAAATTTATCTGATTTGATATTTGAGTATAAA 5751  
Db 7006 AGTTTATTAATATGGAATTTTATATAGAAAAATTTATCTGCTATTGATTTT-AGTATAAG 7064  
Qy 5752 GCAATAATATTTATGATAATAACTATAGAAAAAGATATCTTAGGCTTTTAAATAAACACA 5811  
Db 7065 GCAATAATATTTATGACAATACTATGGAACCAAGATATCTTAGGCTTTTAAATAAACACA 7124  
Qy 5812 TGAATATCATAAATCTTCTGCTTTGTAATTTTCTCCCTTTAATATCAACAATACCATCA 5871  
Db 7125 TGGATATCATAAATCTTCTGCTTTGTAATTTTCTCCCTTTAATATCAACAATACCATCA 7184  
Qy 5872 TCGTCATCATTTACCAATCATTTCTCATGACTTTCATGCTTGCATCATATTTATCTCGTAAAG 5931  
Db 7185 TCATCATCATTTACCAATCATTTCTCATGATTTTCTGCTTGACCATATTTACTGTTAAA 7244  
Qy 5932 TTTG 5935  
Db 7245 GTTG 7248

## RESULT 7

US-09-949-016-17185  
; Sequence 17185, Application US/09949016  
; Patent No. 6812339  
; GENERAL INFORMATION:  
; APPLICANT: VENTER, J. Craig et al.  
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED  
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF  
; FILE REFERENCE: CL001307  
; CURRENT APPLICATION NUMBER: US/09/949,016  
; CURRENT FILING DATE: 2000-04-14  
; PRIOR APPLICATION NUMBER: 60/241,755  
; PRIOR FILING DATE: 2000-10-20  
; PRIOR APPLICATION NUMBER: 60/237,768  
; PRIOR FILING DATE: 2000-10-03  
; PRIOR APPLICATION NUMBER: 60/231,498  
; PRIOR FILING DATE: 2000-09-08  
; NUMBER OF SEQ ID NOS: 207012  
; SOFTWARE: FastSeq for Windows Version 4.0

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; SEQ ID NO 17185
; LENGTH: 8888
; TYPE: DNA
; ORGANISM: Human
US-09-949-016-171

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Query Match 11.0%; Score 651.8; DB 3; Length 8888;  
Best Local Similarity 53.3%; Pred. No. 1.3e-142;  
Matches 3063; Conservative 0; Mismatches 2127; Indels 555; Gaps 58;

Query Match	11.0%;	Score 651.8;	DB 3;	Length 8888;
Best Local Similarity	53.3%;	Pred. No. 1.3e-142;		
Matches 3063;	Conservative 0;	Mismatches 2127;	Indels 555;	Gaps 58;
QY	356	CTCTCCCTCTCAGTTTATCAACTTTTGTACACTTGTGGATCGGTATGCGTGTCCGACGAA	415	
DB	2116	CTCTCCAGTACACAGTTGCTCGAGTTAGATTGTTGCAATGGCCGCGCTTCGAGAA	2175	
QY	416	ATCTATGAGTTTTTCCCTTATATGGGACATTTGGCGCAGCTGCTGCTGTTCTCATTTGCGCT	475	
DB	2176	ATCTGTGAGCTCTTCTCTTATATGGGACCCCTGGCCACCAAGCTGCTCTCTCTTGTGGCGCT	2235	
QY	476	GTGGGCCACGAGGACAAATGGCTGCCCATCAACCCCGGTGCAAGCTCTGAGGTGTCCTCA	535	
DB	2236	CTTGGTACAGGAGGAGAGCTGGCCCATCAGCTCCACATGCAAGGCTTGACAAAGTCCAA	2295	
QY	536	CTTCCAGACGCGTACATCGTCAACCGCACCTTTATGCTGGCCCAAGGAGGTACAGTGC	595	
DB	2296	CTTCCAGACGCTTATATCAACACCGACCTTCATGCTGGCTAAGGAGGTATACATCTC	2355	
QY	596	TCTCTTTCTC-----TCCATACCGCTTGGCAATTTCTCTGAA	632	
DB	2356	AATCCTGCTCTTCTCGTTGGATCTACTTGGAAATCCAAATAGTTCTTAAACTTTTCTTCA	2415	
QY	633	GCATTTGCAAACTCTTTAGGGCGCTTATCTCCGAGGCTCTCACTACCTATGTTTCTG	692	
DB	2416	GAGCATCTCTAAGAGCTTTAGGAACCACTGTTTATCCCTGAGGGTAGATAAATTTCTG	2475	
QY	693	TCTCTTTAGAGACTCTTTTAAGGACTGGATCTTTTCTATTCTTATTTCAAGGTCTCAGGA	752	
DB	2476	TTTTTTTCAGAGACTCTTTGGGAATCTGGCTTTTTTTTTTCTTTGAATCTCTTC-----	2529	
QY	753	CCATTTCTATCTTTGGCGCTTCAGACACATATCTGAAATTTTATCTACAGAGGCGGTTTT	812	
DB	2530	-----CTTCCATTTTGGCGCTTTATGATACATATGATGAATTTTTCCTCCAAAGAGCGGCAAT	2585	
QY	813	AGAAAGCCACCCACGACTGCAATACTTTCC--ATCTGTTGTGCTCTCTTCTCAACTCA	869	
DB	2586	CAGTAATCCATCTGATGAATTTTTTTTTTCCTTTATGCTCTGTCGANTGTTCTTAACTCA	2645	
QY	870	TACTCTCTTGGCTACTC-----CTGAGACCCACTGCGAC	904	
DB	2646	TGCACATCTGAATCTGCTTTTATGATGTTGCTCTGGGAGACGGATGGG	2705	
QY	905	ATACATCTTACTTTACAGGCTTTTCTCCATCTCTTGTACCCAGGCACTTAGGGTTTT	964	
DB	2706	GCATATGTCATGTATAAATTTTTTCTATTTGCTCAAGTCCAGACCCCTTAGTCTTTTT	2765	
QY	965	C-TCCTTTTCAGGCGAGCTTTGCAGATAAACACACAGCTCCGGCTCATCGGGAGAAA	1023	
DB	2766	CTTCTCTTCAGGCTAGCTTGGCTGTATAACACACAGCTTCGTCTCATTTGGGAGAAA	2825	
QY	1024	CTGTTCCGAGGAGTCAAGTGAAGTCTCATCTGTGATGACGAGGGC-----TA	1070	
DB	2826	CTGTTCCAGGAGTCAGTGTAAAGCTACAGTTTGTGACGAAACGGGCGCGTGTCCGCTCCATG	2885	
QY	1071	GCTCGGGAGCTGTGGACCTCTGGGATAG---TCTGAGTATGACCCCTGCTGCTTC	1126	
DB	2886	GGTACTTTGGGGTGTGTGATGATGGTTTAGGTCTTATCCCTTATGACCCCTTCTGTTC	2945	
QY	1127	TTGTCTACCTCAGGCTAAGGATCAGTGTACTCTGATGAACAGGCTGCTCAACTTCCACC	1186	
DB	2946	CTTCCACTGCAGATGAGTGAGCGCTGCTATCTGATGAACAGGCTGCTGAACCTTCAACC	3005	
QY	1187	TGGAAGACATTTCTGCTCCCCAGTCAAGACAGGTTCCGGCCCTCATGACGAGGTGTGTC	1246	
DB	3006	TTGAAGAAGTCTGTTCCCTCAATCTGATAGGTTCCAGCCTTATATGACGAGGTGTGTC	3065	

Qy	1247	CTTTCCTGACAAACTCAGCAATCAGCTCAGCTCCTGTGTAAAGTCTGGCTCTGGCTACCT	1306
Db	3066	CTTTCCTGGCCGGCTCAGCAACAGGCTAAGCACATGTGTAAAGTTACAGCTCTCAGCCTAT	3125
Qy	1307	ATGCTCTCTCTCTTCTTCTCTTCTTATTCACAGTAAGAACCCGAGGTCTGCGCTCTCTCTCT	1366
Db	3126	GCCCACTACCCCTCTCTTCCCTCTCTTCCACAGAGACCCCTTACCCTCACTCTCTCTCTCT	3185
Qy	1367	TCACAAGATGAGGAGGCGCTCAGCACCAACCATCATAGGCACCTTGAATAGGTCAAC	1426
Db	3186	TCCCTTACCCCTAAGCTACGAGGAAGATGCTCTTGGCAGCAGTGTATCAGGAGTCA-	3244
Qy	1427	AAAGCTTTGGCTTCAATGAGTAATACTTTTGTAGTTTGTATATAGTTAAAGCTTTATTTGTT	1486
Db	3245	-----TTTGGGATCATAGAGTATTTTGTCTTTTGTCTTTGCTGCTGAGTCACTCTTGAGTTTA	3298
Qy	1487	TTATCCATGGNAAGAAATCAACTCAATTTCTGTAGGATGAGAAGATGTTGGGAACGAAA	1546
Db	3299	TAGTGTGAATGGGTCTGGAACCTTAAAGTGTACAGAAGCCGCATTTGGTTTGTCTTCCGAA	3358
Qy	1547	AAAGGCTTAGATAGAGAAAACAGATCTCTCAGTAGTACAGTACTTATTTGGGGGGGGGGCAGG	1606
Db	3359	AAAAGGCACTCAGTTGCGTAAAGTAGAAGAAAGGTGTTGGGAAAAAATCTAGATG--TG	3416
Qy	1607	GGGCGATATCCACTAGTCCAAGTACTTTGTTGGGAGAGAAATCOACTGAGTACAAGTACT	1666
Db	3417	GAATGATCCATTTAGTCTAAAGTTGTTGAGGGAGGGGATGGCATGGAGAGAAATTAGA	3476
Qy	1667	TGTGGGGGAAGGAATGCGCACAGAGCAAAAGTTGNAAGGAAAGAGGAAGATGGAGAGCGCT	1726
Db	3477	AGAAAAAGTGGGAAATGGGAAGGCTTAAAGTCG-----GTGGTGGGTGGCGAGACTG	3528
Qy	1727	CAATGTTGGGGGTGTGAAGGTCACCTCTTTTCCATGTGATGAGAGAGTTAAGAAAAATC	1786
Db	3529	TTGGCCTGTTGATGTATGGAAGACCAAAATCGAGGGCGTGTGAACCTTGTATGCCCGCTG	3588
Qy	1787	AGTGTGTGAGTTTGATGCTTTCAGACACCCCAACTATGGCAGACTGTGGGAGACCTCGCA	1846
Db	3589	AACATTTGAACTATGAATAAAAGTTGAGTGGAGTGGGCCCAAGTAAAGGCCCTAGGAC	3648
Qy	1847	TTTAGGA-AGGCGCGCTTTTCACAGAGAACTTTATGCTCATCTCTTGTGCTACACT	1905
Db	3649	TTACTGAAGAGGCTTAATTTTACATGTAGATGTTTTATGTACATTTCTTGTCTTAAGCA	3708
Qy	1906	CCACCTTTGATGAGGTTAAGCTCAGGTTTCGTTCT------AC	1944
Db	3709	TGCAATTTTCTGGAGATACGATTAGGTTTATTTCTTACAGAAATTTGCATAAATCTACTC	3768
Qy	1945	CGTCTTGCTACTGGTGGAAACTTCAGTAGGATTCGCAAGAGACGAGCAGCTCTTCTG	2004
Db	3769	CGCTCTTTCACAAATGCAAACTCAGTAGGATTTCCAAAGATGAAGAGAGGTCTCTTG	3828
Qy	2005	TAAGGAGGACCTTGGATTTTCAGTGTCTTACAGAAACGAAATAGCTCAGAGAACTAGGTC	2064
Db	3829	TAAGGAAGTGACTGGATTCTGGCGTCCAAAGGGAATTCAGAGGCTCAGGAAATCTAGGTC	3888
Qy	2065	AACGTGAATCTAGGTCACAGCGGGCAAAATGACTCAACGCCCTTATTCAGGTGAACG	2124
Db	3889	ACTGTTGAAATCTAGGTCATTTGTGGGCAGAAATTAATAAGAGCTTTAATTCAGGTGAATT	3948
Qy	2125	GTCACTGTCCTCAGATATACTGAGGTATTTGGGCTCCCAACCGGATAAGATCTGTGTAGTA	2184
Db	3949	GTACTGTACCTCCATGGGTGTGAGGTTTCATAAGTTTTCAGCAACATTTAAGATAGTTA	4008
Qy	2185	-GTCTGCTTTTATTTTCAGCACATCAGTGGTGTACGACCAAGAACATCCAGAGAAGTATCA	2243
Db	4009	TGCTTGTTATTTTATATGATATTTGAAGGTGATGACTGCATATCCAGAGGAATGTC	4068
Qy	2244	GAAGGCTGAAGGAGACAGTGAATAAGGTACTATTTGGCAAGCCCAATATCAATAGCCATTCA	2303
Db	4069	AAAAGCTGAAGGACACAGTGAATAAGGTAGGACTGTGATAACTGTCAATGCTAAGTCATGCA	4128



Qy	2304	GTAGGAGACGTGGGATTTCTTTCTCTGCTTCCAGTCT--CTTCTACTTTGTGAACAATTT	2361
Db	4129	ATAGGAGAGACAAATGTTGTTTTTTCTTTCTCTTTTCCCATCACTTTGTGATTTT	4188
Qy	2362	TCCTTTGACTTGTCTACTGTCGTGGTCCAACTACTCACTTAGCTGCACCTGCATCTAGCTGGG	2421
Db	4189	CACTTTGATTTCTCTTACCACAGGCGGATTACT----TTGGTGTCTGTGTATGTAGATATA	4244
Qy	2422	TCTATAGATCTTTCAATCTGTGTCTAAATTT--GTAAGTCACAAATTTCTCGAGCTAGCAG	2478
Db	4245	TCATATATCTAGATGTCAGTTTCCAATCTTGCAATTTGTAGNAATTTCTAGAACTGGTTG	4304
Qy	2479	AAAGCTTAGCTCAGCCAGTCTCATGAGCACTGTGCTCGGAGATGGCTTTGTGACAGAGTCA	2538
Db	4305	GGATCTTAGCTTGTCTAGTCACATAAACCTCAGATTTCTGGGGATGGTCAGTGGCAGAGATA	4364
Qy	2539	ATGCTAGAGACGATCCCTGATTTCCAGCTCTGCAC--TTGCCTAGTGGCCACGTGTAA	2597
Db	4365	GGGCTAGAACTCAGGCTCTCTGAAATCCCAAGCCAGCACTTTTCCGGTGGTGATACAGAT	4424
Qy	2598	TTACTTTAGCCGTAAAGTAATTTGGGAAA--GCCAATTCCCACCGACCTTACATAATCCG	2655
Db	4425	TAGTTTTGGTACCAATTAATTTCTTAGGAAATTTCAGATTTCTATTGACTCATGTATCTG	4484
Qy	2656	AAGAAGCATGCAITTTGAAAACATAGAAA----GCTGGGCACAAACTTACTAGAGATGATTTT	2711
Db	4485	AAGAAGTACTTTGTTTTAAAAACAGAAAAATGCTTATGGCAAAATTTATTTTGAAGTCAITTT	4544
Qy	2712	TGAGCTCATTTAACTGATGCTCTGAAATGTGATCAAAATCAACCAGAGAAATTAACAACAAAG	2771
Db	4545	TGAAGTCATTTAATGCATTTGCTTTTGAACCTTTGGAAGAAATAAAGCTCAGAAACAATGAGAAAG	4604
Qy	2772	AGCTGGATTTTCCAAATAGGACCAAGTATTTTGAATCACTGGTATTAAACAGCTGTCTCTTA	2831
Db	4605	AGCTGGACTTCGATATAGGGCTTAATTTCTGGAGTAA-----TAAACACTTATTTTGA	4656
Qy	2832	ATTTAAATATATAGTCTCTATTTTAGCTGCCTATTTTAAGATTAAACAACAGAGTGGATACCTT	2891
Db	4657	ATTATCATATATCT--ATCAGATATTGATTATAGTTTTAAAAAGCAAGAGACAGACAAC-C	4712
Qy	2892	CCCAATTTACTGGGCGTGGTTTTCAATAGAGTAAAAAATATCAGTCATAGATTAATATAGT	2951
Db	4713	CCGATCTCTTTTATACAGAGTTTCAATAGAGTAAAAAATATTAGTAAGAGATTTATTAAGT	4772
Qy	2952	GTCATGAAAGTATCAGTTTGGAAACCC--TTTCTCTACTTTTATACCTTCATTTCTTAGTT	3008
Db	4773	TAAATGGAAAGTCTGAATTTGGTAAGCTTTTTTTCTCTCTCTCCCAATCAAGACCTTCCA	4832
Qy	3009	ATTATTTTTTTTTCTTCACACCCCTGATCAAGCCACTAGTAAGCACTTATCTGCTGCGAGC	3068
Db	4833	TTCTAGTTTCTTCTTCACTCCCTCAACAAATCCTTAGGGAGCAITTTATCCATGCTGGGC	4892
Qy	3069	TATTATATGACTTTACAGCAAAACAATGCTGTGTGGCCCTCTTTGGGGAAGGGAAACAGG	3128
Db	4893	TGGTGTAATTTCTATAGTGAATGATACCAATCATGTGGCCCTATTTTGGTGGAAAGAAACAAC	4952
Qy	3129	ATACAGAGAGCTCAGGCTAGCAAGTCTGGACTCAACTTAAGCCAGAGGCATGGTTGAT	3188
Db	4953	AATGGA--AGGCTTAGACTATACAATAGT-GACTCACCCCAAAACGGAGGAATGATTAGG	5009
Qy	3189	AGCAGAAAGGTGAGGCTCTTTCAAGAATGGGTGTCTTAAGTAATACAGAAACAGAAAGGC	3248
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Qy	3249	TCTGGTTGATGGAATATACAGTATAGATATCTACCCCTTATCTCC-----TTCTTCTTATA	3301
Db	5069	TCCAGTTGATGGAATTTTTCAGTAAACAAGCTTTAACTTAATTTCCCCCTTTTTTCCCTCTTGA	5128
Qy	3302	GAAGCTAAACCGTCTCTCTCTTCTGTGTGTAGGCTGATAACACAGCTTGTT--TTCTTTTT	3359
Db	5129	CTTTTTAAAAAAGCGTTTCTTCTCTGAGCATCAATTAATGAGTGTGACTGTGTTCTCTCTTT	5188
Qy	3360	GAGTGTTCATGGCTTTTGAGATTTTTTCACTGTCTCGCAGTTCCTTGT--TAGAGGTTTGT	3417

Db	5189	GATAATTGAAGGCTTTGTAGTTTAAATTTGTGAAGCCCAAGTCTCTTGTTATAGAACTAT	5248
Qy	3418	TACCTTGACACCTGGGCTGGATGTAGCATGCCAAGGCACACACTCTCTGAATGCCTGT	3477
Db	5249	TATCTAGACATGAGGGCTGAATGTAGCATGCCACAGCAAGGCATGCTTTACACATCT	5308
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Db	5309	TGCTTAAAAAATTTACTGATTTTCATCTTGCTTGTTGTCTTTAGAAAAAGTGAAGTGTGAGAG	5368
Qy	3530	AGAAAGAACTCAGGAGATGTATTTCTCTGTAGGAAAACTTTTTTTTCCCTTAAAGCC	3589
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Qy	3590	TATAATCCACTTTTCAGTCAACTTTTGACTTTTATACCATGCTGTGCATGAAAGAGTGT	3649
Db	5428	CAATTTTATATTTGCAATGGGTGCCATGTGGAGAGTGAATATGCTTTTGTCTGCTAG	5487
Qy	3650	AGGCCCGCTCTCGTGGCTCTGGGAAAGCACCAATAGGGGAAGAAATGTTATGCCAGAA	3709
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Qy	3710	ATCTGACTGCCAGGAAAACTGGGTCAAGCTCCCCAAAGCCACTACAGGTGTTAAGTAG	3769
Db	5548	CTGTCAAGACTGTCTGAAATAGGGTGGTTTGGGAGGCATTAATCCCTCTCGTGGGGG	5607
Qy	3770	GMACAGTCGAGGTGGTTCATATA--TAATAGAAATGGAACAGAGGGAGGGAAGATAAGCTA	3827
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Qy	3828	CAAAAGTTTCATAGGCTCT--AAGTCTTTAAGATACAAAATAGCTG--GTTGGGCTTCATA	3884
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Db	5728	TCAAAAGAAATATGGAAGG-----TGAAGGGCGGAAGAAAGCAGGGAAGAGAA	5777
Qy	3945	GTAGAGATTTGAAAGACTACAAATCCTCCACAGAGGATTTTTCTTTGGAGGAATCTAGA	4004
Db	5778	GAACCATGTATATA---TAGAGACAATGGTGACAGGTTTTCTTGAAATAT	5829
Qy	4005	ACAAGGTGGTGGATTAGGTGGATCGCAGAAGCACTTGCTTGCCATTTGAAATCTCGGTT	4064
Db	5830	GCAATATGATAGATTAGAGAAATTCAGTAGGGAATGCTTTTCACTTGAAATTTGGGTTT	5889
Qy	4065	TTTGTCCTCCATTGAGTTGAGAGCGTCAACCCTTTTACCTCGATAGGAGGAGGAAA	4124
Db	5890	CCTCT---TCGATTAAGTTTGGGATCCTCATCTGCAATTTGACTTGGAGAG-----AGAAA	5941
Qy	4125	GAAGGGTGTTTTGACTCCTACCTGGAGTTTTACTAGTTTACGCAATGGAACAGACACTC	4184
Db	5942	GAATGAATGTAGGACCTATATCTGGTTTTCTATTAACCTAAGCAAGTGGAA-----	5994
Qy	4185	GGGACCTCCTCTTGACAGAAAAAAGAAAAAGAAACCTGTGTGTTTTCTCTGTGTTG	4244
Db	5995	-----AGACTTATTTGGTATTTTCCCAAAAAGTGAAGAACTTTTCTTTTACTGTTG	6047
Qy	4245	TTCTTTTGTAAAGAAAGCAAGGCAGCTGGGCATGGTGGCCCATGCGTTTAATCCAGCA	4304
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Qy	4305	TTTGGGAGGCAGGCGAGGTGACTTTCTTAATTTCAAGGCCAGCTGCTGTACAAAGTGAG	4364
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Qy	4365	TTCCAGCAGCAGCCAGGGCTATACAGAGAAACCTGTCTCGGGAAGAAAAAAGAGAA	4424
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Db 6202 ACAATAGCTGATGAGCTAAAAAGTCCATGCTATAATCTCATCTGTTTTTAAGAAAGTT 6261  
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Qy 4545 GAAG 4604  
Db 6280 ----- 6279  
Qy 4605 GAAAAAGAAAG 4664  
Db 6280 -----TGCAATTCAGAGCTG 6293  
Qy 4665 ACATGGGACCTATGCTGGTCTTTGAGACAAGGCTTTTGAATTTGAGCGCTCATCAATAGTT 4724  
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Qy 4725 GATCATGGTCAGGTGAGAGGCTACCTGTCAAGCGGAGCCCTGCTGGCTTACGACTTAACA 4784  
Db 6351 TGTCAATAGATAGTTTGGAGCTGCMAAACAGGCCAGGCATAATGGGTGGCACTCGGGAT 6410  
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Db 6411 CCCCCAGATCCCAGCCCTCACTTCAGTCTCTGCTGCTTGGTTAAGAGGGGTGGTCAACTC 6470  
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Db 6471 TCTGCCA----- 6478  
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Db 6479 -----GCTTTTA 6485  
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Db 6486 AACAGCTTCAATAGTGTGAGGTGCACCTGAAATGTAGTGTGTGGGCTCTTCAGTCC 6545  
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Qy 5085 ATGTTGATTAATCAATCTTTTAGAGAAGCGTGATCTTGGTTTTGGTGCTCAGCAAGCAAA 5144  
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Qy 5145 TGTCAACAGCTCTTCTAAGTACCTTACCTTTAGAAAATGCTACCCGTCTCAAAATGGT 5204  
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Qy 5205 TTGTATTCTTATTTTATAGCTTGGAGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGG 5264  
Db 6711 TTGCAATTTATTTTACAGCTTGGAGAGATGGAGAGATCAAAAGCAATTTGGAGAACTGG 6770  
Qy 5265 ACCTGCTGTTTATGCTCTGAGAAATGCTTGCCTGCTGAGCGAGAGAAAGCTTAGAAAAAGA 5324  
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Qy 5325 AGAAGCTGCTTCTGCTTCTTAAAAAGAACAAATAGATCCCTGAATGGAATTTTAA- 5383  
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Qy 5384 ---CTAAGGAAAGTGAAGAGCTAACGTCACCATCAATTAAGAGATTTTCATGAAACCT 5440  
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Qy 5441 GGCTCAGTTGAAGAGAAAAATAGTGTCAA--GTTGTCCATGAGACCAG-AGGTAGACTTG 5497  
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Qy 5498 ATAAACCAAGATTCATTGCAATATTTTATGTCATTGAT-----AATGCAACAGAAAA 5553  
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Qy 5773 AACTATAG-----AAACAAGATATCTTAGCTTTAATAATAACACATGAATATCATAAATCTT 5828  
Db 7311 AATTATAGAGCTATAACATGTTTATTGTGACCTCAATAAACACTTGGATATCCTTAAGTCTT 7370  
Qy 5829 CTGCTTTGTAATTTTCTCCCTTTAATAATCAACAATACCATCATCGTCATCATTTACCCAA 5888  
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Qy 5889 TCATTCTCATGACTTCATGCTTGACTCATATATTATCTGGTAAAGTT 5933  
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RESULT 8

US-09-419-568F-25  
; Sequence 25, Application US/09419568F  
; Patent No. 6331613  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFF) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/419,568F  
; CURRENT FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 25  
; LENGTH: 4797  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-09-419-568F-25

Query Match 11.0%; Score 650; DB 3; Length 4797;  
Best Local Similarity 56.5%; Pred. No. 2.6e-142;  
Matches 1863; Conservative 0; Mismatches 1285; Indels 152; Gaps 29;

Qy 356 CTCTCTCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGGCTGCTCTGCGAGAA 415  
Db 29 CTCTCTCCCAAGTCCACAGTTGCTCGAGTTAGAAATGTCTGCAATGGCCGCCCTGCGAGAA 88  
Qy 416 ATCTATGAGTTTTTCCCTTATGGGACTTTTGGCGCCAGCTGCTGCTTCTCATTTGCCCT 475  
Db 89 ATCTGAGTCTTTCTCTATGGGACCTTGGCCACAGCTGCTCTTCTTCTTGGGCT 148  
Qy 476 GTGGGCCAGAGGCAAAATCGCTGCCCATCAACACCCGGTCAAGCTTGAAGTGTCCAA 535  
Db 149 CTTGGTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGCGAGGCTTGACAAGTCAA 208  
Qy 536 CTTCCAGCAGCGGTACATCGTCAACCGCACCTTTATGCTGGCCAGAGGAGGTACAGCTGCA 595

Db 209 CTTCCAGCAGCCCTATATACCAACCGCACCTTCATGCTGGCTAAGAGGTATACATCTC 268  
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Qy 656 GCTTTATCTCCGAGGTCCTACTACCTATGTTTTCTGTCT-----CTTTAGAG 703  
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Qy 704 ACTCTTTAAGACATGGATCTTTTCTATTTCTATTTCAAGTCTCAGGACCAITTTCTAT 763  
Db 389 TTTTTTCAGAGACTCTTTTGGAAATCTGGCTTTTTTTTTTCTTGAACCTCTTCCTCCAT 448  
Qy 764 CTTGGCTTTCAGGACACATATACCTGAAATTTTATCTACAGAGCGCGTTT--AGAAAGCCA 821  
Db 449 TTTGGCTTTATGATACATATGATGAATTTTCCAAAGAGCGCCATTCAGTAATCCAT 508  
Qy 822 CCCACGACTGCAATCTTCCATCCTGTGTGCTCTCTTTGAACTCATACTCTCTTTGGC 881  
Db 509 CTGATGATTTTTTTTCTTATGCTCTGTGCAATTTGTTCTTAAACTCATGCACACATCTG 568  
Qy 882 TACTC-----CTGAGACCCACTCGGACATACATCTCTAC 916  
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Qy 1259 AACTCAGCAATCAGTCTCAGTCTCTGTGTAAGTCTGGCTACCTATGCTCTCTCTCT 1318  
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Qy 1679 AATGGCACAGAGCAAAAGTTGAAGGGNAAGAGGAAGATGGAGAGCCCTCAATGTGGGGG 1738  
Db 1399 AAATGGGAAGGCTTAAAGTCG-----GTGGTGGGTGGCAGACTGTGTGCCCTGTGTA 1450  
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Qy 1799 TGTATGTTCCAGACACCCCAACTATGGCAGACTGTGGGAGACCTTGGCATTTAGGGA-AGG 1857  
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Qy 1957 TGTGGAAAACTTTCAGTAGGATTTCCCAAAGACGAGGACAGCTCTTCTGTAGGGAGGGAC 2016  
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Db 1811 TAGGTCAATGTGGGCAAAATTAATAAGAGCTTTAAATTCAGGTGAATGTACTGTACTC 1870  
Qy 2137 AGATATATCTAGGTATTGGGCTCCACCGGATAAGATTCTGTGTAGTGA-GTCTCTCTTTTA 2195  
Db 1871 CATGGGTGTGGAGGTTTCATAAAGTTTCAGCAACAATTAAGATAGTTATGCTGTATTG 1930  
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Qy 2316 GGGATTTCTTCTCTGCTCCAGTCTCTTCT--ACTTTGTAAACATTTTCTTTGACTTGT 2373  
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Qy 2374 CTACTGTCTGGTCAATTAATCTACTTAGTGCACCTGCACTAGTGGGTCTATAGATCTT 2433  
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Qy 2434 TCATCTGTCTCTAAATTT---GTAAGTCAAAATTCGAGCTAGCAGAAAGCTTAGCTC 2490  
Db 2167 GATGTGAGTTTCCAAATCTTGCAAAATTTGTAGAAATTCAGAACTGTTGGGATCTTAGCTT 2226  
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Qy 2551 AGCATCCCTGATTTCCAGCTCTGCAC--TTGCCCTAGTGGCCACGCTGTAAATTAATCTTTAGCCT 2609  
Db 2287 AGGTCTCCTGAAATCCCAAGCCAGCACCTTTTCCCGTGTGTATACAGATTAGTTTGGTAC 2346  
Qy 2610 GATTAAGTATTTGGGAAA--GCCAATTTCCACCCAGCTCATATATCCGAGAGACATGCA 2667  
Db 2347 CATTAATTTCTTAGGGAATTTTCAGATTCCTATTGACTCATGTATATCTGAAAGAGTACTTG 2406

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Qy 2668 TTGAAACTAGAAA-----GCTGGGCAAACTTACTAGAGATGATTTTTTGGAGCTCATTA 2723
Db 2407 TTTAAACACAAAATGCTTATGGCAAAATTTTATGAAATCATTTTTTGAAGTCATTA 2466
Qy 2724 ACTGATCCTCGAATCTGATCAATCAACCCGAATAACAAACAAAGAGCTGGATTGC 2783
Db 2467 TGCATTCTTGAACCTTGAAGAAATAAATCTCAGAACATGAGAAAGAGCTGGACTGC 2526
Qy 2784 AAATPAGGACAGTATTTAGAACTCACTGGTATTAACAGCTGTCTATCTTAATTAATAATATAG 2843
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Db 2755 CCTTCACTCCCTCAACAAATCCCTAGGGAGCATTTATCCATGTGGGCTGGTGTACATTT 2814
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Db 2873 TTAGACTTAACTAGT--GACTACCCCAAAACCCGAGGAATGATTAGGACAGTGAAGT 2931
Qy 3201 GAGGCTCTTCAAGTGGGTGTGTTAAGTAATCAGAAAACAGGAAGGCTCTGGTTGATGG 3260
Db 2932 GAGGCTCTT--GCAAGCAGGTACAACTAAATACTCAGAAAACATGAAGGCTCCAGTTGATGG 2990
Qy 3261 AATTATCAGTAAGATCTACCTTATCTCC-----TTCTTCTATAGAGCTTAACCG 3313
Db 2991 AATTTCAGTAACAGCTTAACCTTAATCCCCCTTTTCCCTCTTGACTTTTTTAAAAAA 3050
Qy 3314 TCTCTCTCTTGTGTGTAGCTGATAAACAACGCTGTT--TTCTTTTGTAGTGTTCATGG 3371
Db 3051 GCGTTTCTTCTGAGCATCATTTAATGAGTGTGACTGTTTCTTCTTGTATTAATGGAAG 3110
Qy 3372 CTTTGCAGATTTTTCAGTCTCTGCCAGTTCTTGT--TAGAGGGTTTGTACCTTGAACCC 3429
Db 3111 CTTTGTAGTTTAAATTTGTAAGCCAGTTCTCTTGTATTAGAACTATTATCTAGACATG 3170
Qy 3430 TGGGCTGGATGTAGCATGCAAGGACACACACTTCTGATGCTGTGTAAAGGTTAT 3489
Db 3171 GAGGCTGGAATGTAGCATGCCACAGACAGGCAATGCTTTACACATCTTCTTAAAAAAT 3230
Qy 3490 TATTCAATTTACT-----TTGTCTTTGGAAGGTGAAGTGTGTGAGAAAGAACTCA 3541
Db 3231 TACTGATTTCACTTGTGTTGTCTTTAGAAAAGTGAAGTGTGAGAGAGAGAACTCA 3290
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## RESULT 9

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US-09-354-243B-25
; Sequence 25, Application US/09354243B
; Patent No. 6359117
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Pa
; TITLE OF INVENTION: (TIPI)
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.1
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; CURRENT APPLICATION NUMBER: US/09/354,243B
; CURRENT FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 25
; LENGTH: 4797
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-09-354-243B-25
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Query Match 11.0%; Score 650; DB 3; Length 4797;
Best Local Similarity 56.5%; Pred. No. 2.6e-142;
Matches 1863; Conservative 0; Mismatches 1285; Indels 152; Gaps 29;
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Qy 356 CTCCTCCTCTCAGTATCAACTTTTGACACTTGTGCGATCGGTGATGCTGTCCTGCAAGAA 415
Db 29 CTCCTTCCCAGTCCACAGTTGCTCGAGTTAGAAATGTCTGCAATGGCCGCTTGCAGAA 88
Qy 416 ATCTATGAGTTTTTCCCTTATGGGAGCTTTTGGCCGCGAGCTGCTGCTTCTCATTTGCCCT 475
Db 89 ATCTGTGAGCTCTTTCTTATGGGAGCCCTTGGCCACACAGCTGCTCTCTTCTTGGCCCT 148
Qy 476 GTGGGCCAGGAGGCAAAATGCGCTGCCCATCAACACCCGGTCAAGCTTGAGGTGTCAA 535
Db 149 CTTGGTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCACTGAGGCTTGACAAAGTCAA 208
Qy 536 CTTCCAGCAGCCGTACATCGTCAACCGCACTTTATGCTGGCCAGAGGATACAGTGA 595
Db 209 CTTCCAGCAGCCCTATATACCAACCGCACTTTCATGCTGGCTTAAGAGGATATACATCTC 268
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Qy 704 ACTCTTTAGGAGCTGGATCTTTTCTTCTATTTCAAGGTCTCAGGACCAATTTCTAT 763
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Qy 764 CTTGGGCTTCAGGACACATATACTGAATTTTATCTACAGAGGCGCGTTT--AGAAAGCA 821
Db 449 TTTGGCTTTATGATACATATGATGAATTTTCCCAAAGAGCGCCATTCAGTAATCAT 508
Qy 822 CCACAGCTGCAATATCTTCCATCTCTGTCACCCAGGCACTTAGGGTTTTTCTCTTTGGC 881
Db 509 CTGATGATTTTTTTTCTTATGCTCTGTCATTTGTTCTTAAACTCATGCACACATCTG 568
Qy 882 TACTC-----CTGAGACCACTGGGACATACATCTCTAC 916
Db 569 AATTTCTGCTTTTATGCTTTTGTCTCTGGGAGAGCGGATGGGGCAGCATGTCTAT 628
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Db 629 GTATAAATTTTTTTTCTTATTTGCTCAATGTGCAAGCCCTAGTCTTTTCTTCTTCCAG 688
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Db 689 GCTAGCTTGGCTGATAAACAACAGAGCTTCGTCTCATTTGGGAGAACTGTTCCACGGA 748
Qy 1036 GTCAGTGTAAAGTCTCACTGTGTATGAGCAGGGC-----TAGCTGGGGAGCT 1082
Db 749 GTCAGTGTAAAGTCTCACTGTGTATGAGCAGGGC-----TAGCTGGGGAGCT 808
Qy 1083 GGTGGAGCCCTCTCGGATAG---TCTGAGCTATGACCCCTGCTGCTTCTTGTCTTACCTGC 1138
Db 809 GGTGGTGTATGATGTTTGTAGTCTTATCCCTTATGACCCCTTCTGTTTCCCTTCCACTGC 868
Qy 1139 AGGCTAAGGATCAGTCTACCTGATGAAGCAGGTGCTCAACTTCAACCTCGAAGACATTC 1198
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Db 869 AGATGAGTGAGCGCTGCTATCTGATGAAGCAGGTGCTGAACTTCAACCTTGAAGAGTGC 928  
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Db 929 TGTTCCCTCAATCTGATAGGTTCCAGGCTTATATGCGAGGAGTGGTCCCTTCCTGGCCA 988  
Qy 1259 AACTCAGCAATCAGCTCAGCTCCTGTGTGAAGTCTGGCTCGGCTACCTATGCTCCTCTCT 1318  
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Qy 1319 CTTCTCTCTCTATTCAGTAAGAACCCGAGGTCCTGCCCTCTCTCTCTCTCTCTCTCTCTCT 1378  
Db 1049 CTCCTCCCTCTCTCCAGAGACCCCTTACCCTCAACTCTCTCTCTCTCTCTCTCTCTCTCT 1108  
Qy 1379 GGAGGGCTCAGCACACCACCATCATAGGCCACTTGAATAGTGTCAAAAGGCTTTGGC 1438  
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Qy 1439 TTCAATTGAGTAATCTTGAAGTTGTATAGTTAAGCTTTATTTGTTTATCCATGGAA 1498  
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Qy 1499 AGAAATCAACTCAAAATCTGTAGGATGAGAAAGATTTGGGAAACGAAAAAGGCTTAGAT 1558  
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Qy 1559 AGAGAAACAGATCTGCTGAGTACAGTACTTATGAGGGGGGGGGGAGGGGGCGATATCCA 1618  
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Qy 1858 CGCGCTTTTTCACACGAGAACTTTATGCTCATCTCTTGTGCTACACTCCACCTTTGAT 1917  
Db 1571 GCTTAAATTTTACATGAGATGTTTATGTACATTTCTGTGTTCTAAGCATGCAATTTTCTG 1630  
Qy 1918 GAGGTTAAGCTCAGGTTTCGTTTCT-----ACCGTTCTGTCTAC 1956  
Db 1631 GAGATACGATTTGAGGTTTTTATTCCTTACAGAAATTTGCAATAACTACTCCGCTCTTTCCAC 1690  
Qy 1957 TGGTGGAACCTTCAGTAGGATTTCCCAAGACGAGGACAGCTCTCTCTAAGGAGGGGAC 2016  
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Qy 2017 CTGGAATTTCACTGTCTAGAGAAACAAATAGCTCAGAGAACTTAGGTCAACGCTGAAATCT 2076  
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Qy 2077 AGGTACAGCGGGGCAAAATGATGAACGCTCTATTCAGGTGAAACGGTCAAGTGCCTC 2136  
Db 1811 TAGGTCATTTGGGGCAAAATTAAGAGCTTTAATTCAGGTGAATTTGTACTGTACCTC 1870  
Qy 2137 AGATATCTGAGGTATTTGGGCTCCACCGGATATAGATTCTGTAGTGA-GTCTGCTTTTA 2195  
Db 1871 CATGGGTGTGAGGTTTCATAAAGTTTCAGCACACATTTAAGATGTTATGCTTGTATTG 1930  
Qy 2196 TTTTGCAGCATCATGCTGGTGCAGCACCAAGAACATCCAGAGAATGTCAAGAGGCTGAAGG 2255

Db 1931 TTTTATAGCATATTTGAAGGTGATGACCTGTCATATCCAGAGGAATGTGCAAAAGCTGAAGG 1990  
Qy 2256 AGACAGTGAAGAAAGGTACTATTTCGCAAGCCACAATACTAAGCCATTCACTAGGAGAGCTG 2315  
Db 1991 ACACAGTGAAGAAAGGTAGGACTGATTAAGTCTCAATGCTAAGTCAATGCAATAGGAGAC 2050  
Qy 2316 GGGATTTCTTTCTCTGCTCTCCAGTCTCTTCT--ACTTTGTAAACATTTTCTTTGACTTGT 2373  
Db 2051 AATGTTGTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2110  
Qy 2374 CTACTGTCTGTCATTAATCACTTAGCTGCACCTGCATCTAGCTGGGTCTATAGATCTT 2433  
Db 2111 CTACCACAGGGGGATTT---ACTTTGGTCTGTGTATGTAGATATATCTATATATCTA 2166  
Qy 2434 TCAATCTGCTCTATAAATTT---GTAAGTCAAAATCTCGGAGCTAGCAGAAAGCTTAGCTC 2490  
Db 2167 GATGTCAGTTTCCAAATCTTGGCAATTTGTAGAAATCTAGAACTGGTTGGGATCTTAGCTT 2226  
Qy 2491 AGCCAGTCTCATGAGCACTTGTCTCGGAGATGGCTTGTGACAGAGTCAATGCTTAGAAGAC 2550  
Db 2227 GTCTAGTCACATAACCTCAGATTTCTGGGATGGTCACTGGCAGAGATAGGCTAGAAATGC 2286  
Qy 2551 AGAATCCCTGATTTCCAGCTCTGCAC--TTGCTAGTGGCCACGTTGTAATTTACTTTAGCCT 2609  
Db 2287 AGGTCTCCTGAATCCCAAGCCAGCACTTTTCCCGTGGTGATACAGATTAGTTTGTGTAC 2346  
Qy 2610 GATTAGTATTTGGGAAA--GCCAAATTCACCGACCTCATATATCCGAAAGCAATGCA 2667  
Db 2347 CATTAATCTTTAGGGAATTTTCAGATTCCTATTGACTCATGTAATCTGGAAGAGTACTTG 2406  
Qy 2668 TTGAAAACTAGAAA---GCTGGGCACAAAACCTTACTAGAGATGATTTTTTGAGCTCATATA 2723  
Db 2407 TTTAAAAACAGAAAAATGCGCTATGGGCAATTTATTTTGAAGTCAATTTTGAAGTCAATA 2466  
Qy 2724 ACTGATCTCTGAAATGTGATCAAAATCAACCCAGATAACAAACAAAGAGCTGATTTGC 2783  
Db 2467 TGCATTCCTTTGAACTTGGGAATATAACTCAGAACAAATGAGAAAAAGAGCTGAGCTGC 2526  
Qy 2784 AAATAGCAGCAAGTATTTAGAAATCACTGGTATTAACAGCTGTCACTTAATTTAAATATAG 2843  
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Qy 2844 TGTCTATTTAGTCTGCTTATTTAAGATTAAAAACAGAGTGGATAAATTTCCCAATTTACTG 2903  
Db 2579 TCT---ATCAGATATTGATTATAGTTTAAAAAGCAAGCAGACAAC-CCCGATCTCTTTT 2634  
Qy 2904 GGCTGTGTTTCAATAGAGTAAAAATATCAGTCAATAGATTAAATATAGTGTGATGAAGTA 2963  
Db 2635 ATACAGGTTCAAAATAGAGTAAAAATATTAGTAAGAGATTTATTATAGTTAAATGGAAGTC 2694  
Qy 2964 TGAGTTCGGAACC---CTTTCTTACTTTTACCCTTCAATTTCTTAGTTATTATTATTTT 3020  
Db 2695 TGAATTGGTAGCTTTTCTTCT 2754  
Qy 3021 TCTTTCACACCTGATCAAGCCACTAGTAAGCACCTATCTGCTGCGAGCTATTATATGACT 3080  
Db 2755 CTTTCACTCCCTCAACAAATCCCTAGGAGCATTTATCCATGGTGGCTGGTGTACATTT 2814  
Qy 3081 TTACAGCAAAACAAATTTGCTGTGTGGCTCTTTTGGGGAAGGGAACAGATAGCAGGAGGC 3140  
Db 2815 CTATAGTGAATGATACCATCATGTGGCTATTTTGGTGAAGAAACA--ACAATGGAAGGC 2872  
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Db 2873 TTAGACTAAACAATAGT--GACTCACCCCAAAACCGAGGAATGATTTAGGAGCAGTGAAAGT 2931  
Qy 3201 GAGGCTCTTTCACAAAGTGGGTGCTTAAGTAAATCAGAAAAAGGAGGCTCTGGTTGATGG 3260  
Db 2932 GACGCTCTT--GCAAGCAGGTACAACATAAATACTCAGAAAAACATGAAGGCTCCAGCTGTGATG 2990  
Qy 3261 AATTATCAGTAAGATATCTACCTTATCTCC-----TTCTTCTATAGAGCTTAAACCG 3313  
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Db      928  TACITTTATATGTAAGTTTATTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 987
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Qy      5761 ATTATGATAAATACTATAGAAACAAGATATCTTAGGCTTTTAAATAAACACATGAATATCA 5820
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Db      1108 TAAA 1111

RESULT 12
US-09-354-243B-9
; Sequence 9, Application US/09354243B
; Patent No. 6359117
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Pe
; TITLE OF INVENTION: (Tifs)
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.1
; CURRENT APPLICATION NUMBER: US/09/354,243B
; CURRENT FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 9
; LENGTH: 1111
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-09-354-243B-9

Query Match          10.1%; Score 602.4; DB 3; Length 1111;
Best Local Similarity 99.8%; Pred. No. 2e-131;
Matches 603; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Db      508  AAAGCTTGGAGAGCGGAGAGATCAAAGCGATCGGGAACTGGACCTGCTGTTTATGTC 567
Qy      5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAAAGAACTGCTCCTTCCT 5340
Db      568  TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAAAGAACTGCTCCTTCCT 627
Qy      5341 GCCTTCTAAAAAGAACAAATAAGATCCCTGAAATGGAATTTTTTACTTAAAGGAAAGTGAGAA 5400
Db      628  GCCTTCTAAAAAGAACAAATAAGATCCCTGAAATGGAATTTTTTACTTAAAGGAAAGTGAGAA 687
Qy      5401 GCTAACGTCACACATCATTAGAAAGATTTCATGAAACCTGGCTCAGTTGAAAGAGAAAA 5460
Db      688  GCTAACGTCACACATCATTAGAAAGATTTCATGAAACCTGGCTCAGTTGAAAGAGAAAA 747
Qy      5461 TAGTGTCAAGTTGTCCATGAGACAGAGGTAGACTTGTATACCAACAAAGATTCATTGACA 5520
Db      748  TAGTGTCAAGTTGTCCATGAGACAGAGGTAGACTTGTATACCAACAAAGATTCATTGACA 807
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Db      808  ATATTTTATTTGTCATTTGATAATGCAACAGAAAAAGATGTATGTAATTTTAAAAAATTTGTTTGA 867
Qy      5581 AGGAGGTTTACCTCTCATTCCTCTAGAGAAAGCCATGTAACTTCATTTCCATAACCAA 5640
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Db      928  TACITTTATATGTAAGTTTATTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 987
Qy      5701 ATATGGATTTTATTTATAGAAAAATTTATCTGATGTTTGATATTTTGAGTATAAGCAAAATAAT 5760
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Qy      5821 TAAA 5824
Db      1108 TAAA 1111

RESULT 13
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; Sequence 7, Application US/09178973B
; Patent No. 6274710
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Louhed, Jamila
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: (Tifs)
; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543
; CURRENT APPLICATION NUMBER: US/09/178,973B
; CURRENT FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 17
; SEQ ID NO 7
; LENGTH: 1119
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-09-178-973B-7

Query Match          9.4%; Score 555.2; DB 3; Length 1119;
Best Local Similarity 96.0%; Pred. No. 2.5e-120;
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;

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Qy	5701	ATATGGATTATTTATAGAAAAATTATCTGATGTTTGATATTTGAGTATAAAGCAATAAT	5760
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Qy	5821	TAAA	5824
Db	1109	TAAA	1112

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GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:16 ; Search time 2968.2 Seconds  
(without alignments)  
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Title: US-09-751-797-29

Perfect score: 5935

Sequence: 1 gaattcaagtcacatgcaa.....atatatctggttaagtgtg 5935

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 9793542 seqs, 4134689005 residues

Total number of hits satisfying chosen parameters: 19587084

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Published Applications NA Main:\*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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1	5935	100.0	5935	3	US-09-751-797-29
2	5935	100.0	5935	7	US-10-627-273-29
3	4245.2	71.5	7445	3	US-09-751-797-8
4	4245.2	71.5	7445	7	US-10-627-273-8
5	650	11.0	4797	3	US-09-751-797-25
6	650	11.0	4797	7	US-10-627-273-25
7	602.4	10.1	1111	3	US-09-751-797-9
8	602.4	10.1	1111	7	US-10-627-273-9
9	555.2	9.4	1119	3	US-09-751-797-7
10	555.2	9.4	1119	7	US-10-627-273-7
11	555.2	9.4	1166	5	US-10-084-298-3
12	555.2	9.4	1166	8	US-10-256-977-3
13	555.2	9.4	1166	6	US-10-873-972-3
14	555.2	9.4	1166	10	US-11-157-387-3
15	541.4	9.1	1050	5	US-10-090-365-40
16	541.4	9.1	1050	5	US-10-104-919-42
17	541.4	9.1	1050	8	US-10-807-837-10
18	541.4	9.1	1050	9	US-10-968-432-42
19	541.4	9.1	1050	10	US-11-045-944-40
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21	264.4	4.5	778	7	US-10-395-741B-37
22	264.4	4.5	778	7	US-10-806-294-37
23	218.4	3.7	1177	10	US-11-013-920-1

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25	218.4	3.7	1191	5	US-10-084-298-1	Sequence 1, Appli
26	218.4	3.7	1191	6	US-10-256-977-1	Sequence 1, Appli
27	218.4	3.7	1191	8	US-10-873-972-1	Sequence 1, Appli
28	218.4	3.7	1191	10	US-11-157-387-1	Sequence 1, Appli
29	216.4	3.6	1116	3	US-09-728-911-14	Sequence 14, Appli
30	216.4	3.6	1116	3	US-09-525-055D-7	Sequence 7, Appli
31	216.4	3.6	1116	3	US-09-746-375-1	Sequence 1, Appli
32	216.4	3.6	1116	5	US-10-090-365-14	Sequence 14, Appli
33	216.4	3.6	1116	5	US-10-104-919-14	Sequence 14, Appli
34	216.4	3.6	1116	7	US-10-395-741B-1	Sequence 1, Appli
35	216.4	3.6	1116	7	US-10-806-294-1	Sequence 1, Appli
36	216.4	3.6	1116	8	US-10-807-837-5	Sequence 5, Appli
37	216.4	3.6	1116	9	US-10-968-432-14	Sequence 14, Appli
38	216.4	3.6	1116	9	US-10-981-998-14	Sequence 14, Appli
39	216.4	3.6	1116	10	US-11-045-944-14	Sequence 14, Appli
40	215	3.6	1152	3	US-09-870-574-1	Sequence 1, Appli
41	215	3.6	1152	5	US-10-006-867-153	Sequence 153, App
42	215	3.6	1152	5	US-10-066-500-125	Sequence 125, App
43	215	3.6	1152	5	US-10-063-547-153	Sequence 153, App
44	215	3.6	1152	5	US-10-063-551-153	Sequence 153, App
45	215	3.6	1152	5	US-10-063-616-153	Sequence 153, App

#### ALIGNMENTS

##### RESULT 1

US-09-751-797-29  
; Sequence 29, Application US/09751797  
; Patent No. US20010024652A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; FILE REFERENCE: (TIFS) The Proteins Encoded, and Uses Thereof  
; CURRENT APPLICATION NUMBER: US/09/751,797  
; CURRENT FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 29  
; LENGTH: 5935  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
US-09-751-797-29

Query Match 100.0%; Score 5935; DB 3; Length 5935;  
Best Local Similarity 100.0%; Pred. No. 0;  
Matches 5935; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	GAATTCAAAGTCCACATCGAATCAATCGAATCACTTTGTAATAATCTCTTCTTCAAAATATCC	60
Db	1	GAATTCAGTCCACATCGAATCAATCGAATCACTTTGTAATAATCTCTTCTTCAAAATATCC	60
Qy	61	ATCTATATAGTATAAGTTATTTAGGATCAATTTAAATAATGTTTGTGAGACTTATGTTT	120
Db	61	ATCTATATAGTATAAGTTATTTAGGATCAATTTAAATAATGTTTGTGAGACTTATGTTT	120
Qy	121	GCACAGTAAAGTCCAGAGAAATAGCAAAATGATATAGTATTTTATTTTAAAAAAT	180
Db	121	GCACAGTAAAGTCCAGAGAAATAGCAAAATGATATAGTATTTTATTTTAAAAAAT	180
Qy	181	CTATGCTTAAATGCTTATTAGATTGTTCACTACTGACATTTCCAAACTTTAACTTGACCT	240
Db	181	CTATGCTTAAATGCTTATTAGATTGTTCACTACTGACATTTCCAAACTTTAACTTGACCT	240
Qy	241	TGGCTATGATTTCAACCTTTGTTATTTGTCATCTACCATTAAGTGTGCTCATTACCANGC	300

Db 241 TGGCTATGATTTCAACCTTTGTATTTGCACTACCAATACTGTGTGCTCACTTACCATGC 300  
Qy 301 TATCCGACGAGCATGTTCCTCCCTGATGTTTTTGGCCCTTTTGGCTCTCTCGCTAACAGGCTCTC 360  
Db 301 TATCCGACGAGCATGTTCCTCCCTGATGTTTTTGGCCCTTTTGGCTCTCTCGCTAACAGGCTCTC 360  
Qy 361 CTCTCAGTTATCAACTTTTGGACCTTGTGCGATCGGTGATGGCTGTCTCTCGAAGAACTTA 420  
Db 361 CTCTCAGTTATCAACTTTTGGACCTTGTGCGATCGGTGATGGCTGTCTCTCGAAGAACTTA 420  
Qy 421 TGAGTTTTTCCCTTATGGGGACTTTTGGCCGCCAGCTGCTGCTTCTCAATTGGCCCTGTGGG 480  
Db 421 TGAGTTTTTCCCTTATGGGGACTTTTGGCCGCCAGCTGCTGCTTCTCAATTGGCCCTGTGGG 480  
Qy 481 CCCAGGAGGCAAAATGCGCTGCCATCAACACCCCGGTGCAAGCTTTGAGGTGCCAACTTCC 540  
Db 481 CCCAGGAGGCAAAATGCGCTGCCATCAACACCCCGGTGCAAGCTTTGAGGTGCCAACTTCC 540  
Qy 541 AGCAGCCGTACATCGTCAACCGGACCTTTATGCTGGCCAGGAGGTACAGCTGCATCTCT 600  
Db 541 AGCAGCCGTACATCGTCAACCGGACCTTTATGCTGGCCAGGAGGTACAGCTGCATCTCT 600  
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Db 601 TTCTCTCCATACCGCTTGGCCATTTCTCTGAAGCACTTGCAAACTCTTTAGGGGCGCTTT 660  
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Db 661 ATCTCCGAGGTCTCACTACTATGTTTTCTGTTCTTTTAGAGACTCTTTTAAGGACTGGA 720  
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Db 721 TCTTTTCTATTTCTATTTCAAGGTCTCAGACCAATTCCTATCTTGGCTTTCAGGACAC 780  
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Db 781 ATATACTGAAATTTTATCTACAGAGCGGTTTAGAAGGCCAACCCACGACTGCAATACTTT 840  
Qy 841 CCATCCTGTGTGCTCTCTCTGAACTCATACTCTCTTGGCTACTCTCTGAGACCCACTGC 900  
Db 841 CCATCCTGTGTGCTCTCTCTGAACTCATACTCTCTTGGCTACTCTCTGAGACCCACTGC 900  
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Db 901 GGACATACATCTCTACTACAGGCTTTTCTTCCATCTCTCTTGTCAACCAGGCACTTAGGG 960  
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Db 1081 CTGTTGACCCCTCTGGGATAGTCTGACGTATGACCCCTGCTGCTTCTTGTCTACTGTCAG 1140  
Qy 1141 GCTAAGGATCAGTGTCTCTGATGAAGCAGGTGCTCAACTTTCACCTCGAAGACATTTCTG 1200  
Db 1141 GCTAAGGATCAGTGTCTCTGATGAAGCAGGTGCTCAACTTTCACCTCGAAGACATTTCTG 1200  
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Db 1261 CTCAGCAATCAGCTCAGCTCTCTGTGTAGTCTGGCTCTGGCTACCTATGCTCCTCTCTCT 1320  
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Db 1321 TCCTCTCTTATTTCCAGTAAAGAACCCGAGGTCTGCGCTCTCTCTCTTTCACAAGAGTGAAG 1380  
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Db 1741 TGAAGGTCACTCTCTTTTCCATGTGATGAGAGTTAAGAAATCAAGTGTGTCAGTTTG 1800  
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Db 1801 ATGTCTTTCAGACACCCCAACTATGCGAGACTGTGGGAGACCTGCGCATTTAGGGAAGGCGC 1860  
Qy 1861 GGCTTTTTCACACGAGAAACTTTATGCTCATCTCTGTGTACACTCCACCTTTGATGAG 1920  
Db 1861 GGCTTTTTCACACGAGAAACTTTATGCTCATCTCTGTGTACACTCCACCTTTGATGAG 1920  
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Db 1921 GTTAAGCTCAGGTTTCGTTTCTACCGTTCTGTCTACTGTTGGAACCTTCACTAGGATGCC 1980  
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Db 1981 CCAAAGACGAGGACAGCTCTTCTGTAAGGGAGGACCTTGGATTTCAAGTGTCTTAGAGAAC 2040  
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Db 2041 GAAATAGCTCAGAGAAATCTAGGTCAACGTGAAATCTAGGTCAACGCGGGCAAAAATGACT 2100  
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Db 2161 CACCGGATAAGATTTCTGTTAGTGTGCTGCTTTTATTTTGCAGCACATCAGTGTGACGA 2220  
Qy 2221 CCAGAACATCCAGAAAGATGTGAGAGGCTGAAAGGAGACAGTGAAGGATCTAATTGGC 2280  
Db 2221 CCAGAACATCCAGAAAGATGTGAGAGGCTGAAAGGAGACAGTGAAGGATCTAATTGGC 2280  
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Db 2281 AAGCCACATCTAAGCCATTTCACTAGGAGACGTTGGGATTTCTTCTCTGTTCCCACT 2340  
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Db 2341 CTCTTCTACTTTGTAAACATTTTCTTGTACTTGTCTACTGTCTGGTCCATTTACTCACTTAG 2400  
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Db 2401 CTGCACTGCTACTAGCTGGGTCTATAGATCTTTCAATCTGTGTCTAAATTTGTAAAGTCA 2460

Qy	2461	CAATTC	TGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTCATGAGCACTTGCCTCGGAGGA	2520
Db	2461	CAATTC	TGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTCATGAGCACTTGCCTCGGAGGA	2520
Qy	2521	TGGCTT	GTGACAGAGTCAATGCTAGAGACAGCATTCCCTGATTTCCAGCTCTGCACCTTGC	2580
Db	2521	TGGCTT	GTGACAGAGTCAATGCTAGAGACAGCATTCCCTGATTTCCAGCTCTGCACCTTGC	2580
Qy	2581	CTAGTGC	CCACGTCGTAATTACTTTAGCGCTGATTAAAGTATTTGGGAAAGCCAAATTTCCCA	2640
Db	2581	CTAGTGC	CCACGTCGTAATTACTTTAGCGCTGATTAAAGTATTTGGGAAAGCCAAATTTCCCA	2640
Qy	2641	GACCTA	CATAATCCGAAGAACATGCAATGAAACCTAGAAAGCTGGGCACAAACTTACTA	2700
Db	2641	GACCTA	CATAATCCGAAGAACATGCAATGAAACCTAGAAAGCTGGGCACAAACTTACTA	2700
Qy	2701	GAGATG	ATTTTGTAGCTCATTTAAACTGATGCTCTGAAATGTGATCAAAATCAACCCAGAA	2760
Db	2701	GAGATG	ATTTTGTAGCTCATTTAAACTGATGCTCTGAAATGTGATCAAAATCAACCCAGAA	2760
Qy	2761	AACAA	CAAAAGAGCTGGATTTGCAATAGACAAAGTATTTAGAAATCACTGATTAACAG	2820
Db	2761	AACAA	CAAAAGAGCTGGATTTGCAATAGACAAAGTATTTAGAAATCACTGATTAACAG	2820
Qy	2821	CTGTCA	TCTTAAATTAATAGTGTCTATTTAGCTGCCTATTTAAGATTAACACAAGA	2880
Db	2821	CTGTCA	TCTTAAATTAATAGTGTCTATTTAGCTGCCTATTTAAGATTAACACAAGA	2880
Qy	2881	GTGGAT	AACTTCCCAATTTACTGGGCTCTGTTTCAATAGAGTAAATATCAGTCATAGA	2940
Db	2881	GTGGAT	AACTTCCCAATTTACTGGGCTCTGTTTCAATAGAGTAAATATCAGTCATAGA	2940
Qy	2941	TTAATTA	TAGTGTCATCAAAAGTAGTGTGGAACCCCTTTCCCTTACCTTTTACCTTCAT	3000
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Db	3001	TCTTAG	TATATTTTTTTTTTTTCTTCAACCTCGATCAAGCCACTAGTAAAGCACTATCTG	3060
Qy	3061	CTCGAG	CTATATATGACTTTACAGCAAAACAATTCGTGTGTGGCTCTTTGGGGAAG	3120
Db	3061	CTCGAG	CTATATATGACTTTACAGCAAAACAATTCGTGTGTGGCTCTTTGGGGAAG	3120
Qy	3121	GGACAG	ATAGCAGGAGGCTCAGGCTAGCAAGTCTGGAATCAACCTAAAGCCAGAGGCA	3180
Db	3121	GGACAG	ATAGCAGGAGGCTCAGGCTAGCAAGTCTGGAATCAACCTAAAGCCAGAGGCA	3180
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Db	3181	TGGTTG	ATAGCAGAAAGTGAGGCTCTTCAAGTGGGTGCTTAAGTAATCAGAAAC	3240
Qy	3241	AGGAAG	GCCTCTGGTTGATGGAAATATCAGTAAGATATCTACCCCTTATCTCCTTCTCTAT	3300
Db	3241	AGGAAG	GCCTCTGGTTGATGGAAATATCAGTAAGATATCTACCCCTTATCTCCTTCTCTAT	3300
Qy	3301	AGAAGC	TAAACCGTCTCTCTTCTTGTGTAGGCTGATAAACACGCTTGTTTTCTTTTG	3360
Db	3301	AGAAGC	TAAACCGTCTCTCTCTTCTTGTGTAGGCTGATAAACACGCTTGTTTTCTTTTG	3360
Qy	3361	AGTGTC	ATTCAGTGTGAGATTTTTTCAGTGTCTGSCCAGTCTTGTGTAGAGGGTTGTTAC	3420
Db	3361	AGTGTC	ATTCAGTGTGAGATTTTTTCAGTGTCTGSCCAGTCTTGTGTAGAGGGTTGTTAC	3420
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Db	3421	CTTGAC	ACCTGGGCTTGGATGTTAGCATGCCAAAGGCACACATCTCTGAATGCTGTGTA	3480
Qy	3481	AAAGGT	TATTATTCATTTCTTTTGTGTAAGGGTGAAGTGTGTGTGAGAAAGAACTC	3540
Db	3481	AAAGGT	TATTATTCATTTCTTTTGTGTAAGGGTGAAGTGTGTGTGAGAAAGAACTC	3540

Qy	3541	ACAGGAGATGTAATCTCTGTAGGAAACCTTTTTTTTCCCTTAAAAAGCCTATATATCCACT	3600
Db	3541	ACAGGAGATGTAATCTCTGTAGGAAACCTTTTTTTTCCCTTAAAAAGCCTATATATCCACT	3600
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Db	3601	TTCAAGTCAACTTTTGACTTTTTATACCATGCTGTACATGAAGAGTGTTTTAGGCCCGCTCT	3660
Qy	3661	CGTGGCTCTGGGAAAAAGCACCAATAGGGGAAGAAATGTTATGCGGAAAAATCTGACTGGC	3720
Db	3661	CGTGGCTCTGGGAAAAAGCACCAATAGGGGAAGAAATGTTATGCGGAAAAATCTGACTGGC	3720
Qy	3721	AGGGAACCTGGGTCTAGAGCTCCCCAAGACCACTACAGTGTGTTAAGTAGGAAACAGTCGAG	3780
Db	3721	AGGGAACCTGGGTCTAGAGCTCCCCAAGACCACTACAGTGTGTTAAGTAGGAAACAGTCGAG	3780
Qy	3781	GGTGGGTTCATATAATAGATGNAACAGAGGGAGGNAAGTAAGCTACAAGTTTTCATAG	3840
Db	3781	GGTGGGTTCATATAATAGATGNAACAGAGGGAGGNAAGTAAGCTACAAGTTTTCATAG	3840
Qy	3841	GGTCCTAAGTCTTTAAGATACAAAATAGCTGGTGGGCTTCATAAACAAGGAAGCTCTGGG	3900
Db	3841	GGTCCTAAGTCTTTAAGATACAAAATAGCTGGTGGGCTTCATAAACAAGGAAGCTCTGGG	3900
Qy	3901	AAGGCAGCAAGCANTGAGAGGGAGATGGAAGGGAAAAAACAATGTAGAGGATTTGAAAA	3960
Db	3901	AAGGCAGCAAGCANTGAGAGGGAGATGGAAGGGAAAAAACAATGTAGAGGATTTGAAAA	3960
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Db	3961	GCTACAAATCTCCACGAGAGGATTTTTCTTTGAGGNACTCAGAACAAAGGTGGTGGATT	4020
Qy	4021	AGTGGATCCGAAAGGACTTGCTTGCCACTTTTGAATCTGGGTTTTTGTCTCTCCATTTGA	4080
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Qy	4081	GGTTGAGAGCGTCAACCCCTTTTTTACCTGGATAGGAGGAGAAAGAGGGGTGTTTTGAC	4140
Db	4081	GGTTGAGAGCGTCAACCCCTTTTTTACCTGGATAGGAGGAGAAAGAGGGGTGTTTTGAC	4140
Qy	4141	TCCTACCTGAGTTTTACTAGTTTACGCAATGNAACAGACACTCGGACCTCTCTTTGAC	4200
Db	4141	TCCTACCTGAGTTTTACTAGTTTACGCAATGNAACAGACACTCGGACCTCTCTTTGAC	4200
Qy	4201	AAGAAAAAAGAAAAAGGAAACCTGTGTTTTCTCTGTTTTGTTCTTTTTTTGTTTGAAGAA	4260
Db	4201	AAGAAAAAAGAAAAAGGAAACCTGTGTTTTCTCTGTTTTGTTCTTTTTTTGTTTGAAGAA	4260
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Db	4261	GCACAGCAGCTGGGCATGTGTGCCCATGCTTTTAATCCACGATTTGGGAGGCAGAGGC	4320
Qy	4321	AGGTGACTTTCTAAATTCAAAGGCAGCGCTGGTCTACAAAGTAGTTCCAGGACAGCCAGG	4380
Db	4321	AGGTGACTTTCTAAATTCAAAGGCAGCGCTGGTCTACAAAGTAGTTCCAGGACAGCCAGG	4380
Qy	4381	GCTATACAGAGAAACCTGTCTCGGGAAAAAAGAAAAAGAAAGAAAGAAAGAAAGAAAG	4440
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Qy	4441	AGAAGAGGAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG	4500
Db	4441	AGAAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG	4500
Qy	4501	AGGAGGAGGAG	4560
Db	4501	AGGAGGAGGAG	4560
Qy	4561	AAGCAGAA	4620
Db	4561	AAGCAGAA	4620
Qy	4621	AGAAAGAAAAAGCAGCAAGCACTGGCAATGGCAATGGCAATGGCAATGGCAATGGCAAT	4680



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Qy |||||  
481 CCAGAGGCAAAATGCGCTGCCCAATCAACACCGGGTGCAAGCTTGAGGTGTCAACTTCC 540  
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1501 AAATCAACTCAATCTCTAGGATGAAAAGATGTTGGAAACGAAAAAGCCCTAGATAG 1560  
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1561 AGAAACAGATCTGCTGAGTACAGTACTTATGGGGGGGGGGGAGGGGGGATATCCACT 1620  
Db |||||  
1561 AGAAACAGATCTGCTGAGTACAGTACTTATGGGGGGGGGGGAGGGGGGATATCCACT 1620  
Qy |||||  
1621 GAGTCCAACTGCTGTTGGGAGAGAAATCCACTGAGTACAAAGTACTTGTGGGGGAGGAA 1680  
Db |||||  
1621 GAGTCCAACTGCTGTTGGGAGAGAAATCCACTGAGTACAAAGTACTTGTGGGGGAGGAA 1680  
Qy |||||  
1681 TGSCACAGAGCAAAAGTTGAAAGGAAAGAGATGGAGAGCCCTCAATGTTGGGGGTG 1740  
Db |||||  
1681 TGSCACAGAGCAAAAGTTGAAAGGAAAGAGATGGAGAGCCCTCAATGTTGGGGGTG 1740  
Qy |||||  
1741 TGAAAGGTCACTCCCTTTTCCATGTGATGGAGAGTTTAAAGAAATCAGTGTGTGAGTTG 1800  
Db |||||  
1741 TGAAAGGTCACTCCCTTTTCCATGTGATGGAGAGTTTAAAGAAATCAGTGTGTGAGTTG 1800  
Qy |||||  
1801 ATGCTCTTCAGACACCCCAACTATGGCAGACTGTGGGAGACCTGGCATTTAGGGAAGCGC 1860  
Db |||||  
1801 ATGCTCTTCAGACACCCCAACTATGGCAGACTGTGGGAGACCTGGCATTTAGGGAAGCGC 1860  
Qy |||||  
1861 GGCTTTTCAACGAGAAACTTTATGCTCATCTCTGTGTACACTCCCACTTTGATGAG 1920  
Db |||||  
1861 GGCTTTTCAACGAGAAACTTTATGCTCATCTCTGTGTACACTCCCACTTTGATGAG 1920  
Qy |||||  
1921 GTTAAAGTCAAGTTTCTGTTTCTACCGTTCTTGTCTACTGTGGGAACTTCAGTAGGATTC 1980  
Db |||||  
1921 GTTAAAGTCAAGTTTCTGTTTCTACCGTTCTTGTCTACTGTGGGAACTTCAGTAGGATTC 1980  
Qy |||||  
1981 CCAAGACAGGAGACAGCTCTCTGTAAAGGAGGAGCTGGATTTCACTGTCCTTAGAGAAC 2040  
Db |||||  
1981 CCAAGACAGGAGACAGCTCTCTGTAAAGGAGGAGCTGGATTTCACTGTCCTTAGAGAAC 2040  
Qy |||||  
2041 GAAATAGCTCAGAGAACTTAGGTCAAACGTGAAATCTAGGTCAACAGCGGGCAAAATGACT 2100  
Db |||||  
2041 GAAATAGCTCAGAGAACTTAGGTCAAACGTGAAATCTAGGTCAACAGCGGGCAAAATGACT 2100  
Qy |||||  
2101 GAAAGCTCTATTCAGAGTGAAAGGCTCACTGCTCAGATATATCTAGGATTTGGGGTCC 2160  
Db |||||  
2101 GAAAGCTCTATTCAGAGTGAAAGGCTCACTGCTCAGATATATCTAGGATTTGGGGTCC 2160  
Qy |||||  
2161 CACCGATAGATTTCTGTAGTGTGCTGCTTTTATTTTGCAGACATCAGTGTGACGA 2220  
Db |||||  
2161 CACCGATAGATTTCTGTAGTGTGCTGCTTTTATTTTGCAGACATCAGTGTGACGA 2220  
Qy |||||  
2221 CCAGAACATCCAGAGAAATGTCAAGAGGCTGAAAGGAGACAGTGAAAGAGTACTATTGGC 2280  
Db |||||  
2221 CCAGAACATCCAGAGAAATGTCAAGAGGCTGAAAGGAGACAGTGAAAGAGTACTATTGGC 2280  
Qy |||||  
2281 AGCCCAATATCTAAGCCATTCAGTAGGAGACGTGGGGATTTCTTCTCTGCTTCCAGT 2340  
Db |||||  
2281 AGCCCAATATCTAAGCCATTCAGTAGGAGACGTGGGGATTTCTTCTCTGCTTCCAGT 2340  
Qy |||||  
2341 CTCCTCTACTTTGTAAACATTTCTTTGACTGTGTCTGTCTGCTTCAATCTGCTTAAAGTCA 2400  
Db |||||  
2341 CTCCTCTACTTTGTAAACATTTCTTTGACTGTGTCTGTCTGTCTTCAATCTGCTTAAAGTCA 2400  
Qy |||||  
2401 CTGCACCTGCTAGCTGGGTCTATAGATCTTTTCAATCTGTCTGTCTTAAATTTGTAAGTCA 2460  
Db |||||  
2401 CTGCACCTGCTAGCTGGGTCTATAGATCTTTTCAATCTGTCTGTCTTAAATTTGTAAGTCA 2460  
Qy |||||  
2461 CAATCTGAGGTAGCAGAAAGCTTAGCTCAGCAGCTCTCATAGAGCACTTGTCTGGAGGA 2520  
Db |||||  
2461 CAATCTGAGGTAGCAGAAAGCTTAGCTCAGCAGCTCTCATAGAGCACTTGTCTGGAGGA 2520  
Qy |||||  
2521 TGGCTTTGTACAGAGTCAATGCTAGAGAGCAGCATCCCTGATTCAGGCTCTGCATTCG 2580  
Db |||||  
2521 TGGCTTTGTACAGAGTCAATGCTAGAGAGCAGCATCCCTGATTCAGGCTCTGCATTCG 2580  
Qy |||||  
2581 CTAGTGGCCACGTGTAAATTAATTTAGCTGATTAAGTATTTTGGGAAAGCCAAATTCACCAC 2640

Db	2581	CTAGTGGCCACGCTGAATTAATCTTTAGCCTGATTAAGTAATTTGGGAAAGCCAAATTTCCACACC	2640
Qy	2641	GACCTACATAATCCGAAGAAGCATGCAATGAAAACTAGAAAGCTGGGCACAAAACCTTACTTA	2700
Db	2641	GACCTACATAATCCGAAGAAGCATGCAATGAAAACTAGAAAGCTGGGCACAAAACCTTACTTA	2700
Qy	2701	GAGATGATTTTTGAGCTCATTTAAACTGATGCTCTGAAATGTGATCAAAATCAACCCAGAAT	2760
Db	2701	GAGATGATTTTTGAGCTCATTTAAACTGATGCTCTGAAATGTGATCAAAATCAACCCAGAAT	2760
Qy	2761	AACAAACAAAGAGCTGGATTTGCAAAATAGACAAGTATTTAGAAATCACTGGTATTAAACAG	2820
Db	2761	AACAAACAAAGAGCTGGATTTGCAAAATAGACAAGTATTTAGAAATCACTGGTATTAAACAG	2820
Qy	2821	CTGTCATCTTAAATTAATAATAGTGTCTATTTAGCTGCCATTTTAAAGATTAAACACAAGA	2880
Db	2821	CTGTCATCTTAAATTAATAATAGTGTCTATTTAGCTGCCATTTTAAAGATTAAACACAAGA	2880
Qy	2881	GTGGATAACTTTCCCAATTTTACTGGGCCCTGGTTTCAATAGAGTAAATAATATCAGTCAATAGA	2940
Db	2881	GTGGATAACTTTCCCAATTTTACTGGGCCCTGGTTTCAATAGAGTAAATAATATCAGTCAATAGA	2940
Qy	2941	TTAATTATAGTGTCAATGAAAGTATGAGTTGGAAACCCCTTTCCCTTACCTTTTACCTTCATT	3000
Db	2941	TTAATTATAGTGTCAATGAAAGTATGAGTTGGAAACCCCTTTCCCTTACCTTTTACCTTCATT	3000
Qy	3001	TCCTTAGTATTATTTTTTTTCTTCTCACACCTGATCAAGCCACTAGTAAGCACCTATCTG	3060
Db	3001	TCCTTAGTATTATTTTTTTTCTTCTCACACCTGATCAAGCCACTAGTAAGCACCTATCTG	3060
Qy	3061	CTGCGAGCTATTATGACTTTTACAGCAAAACAAATTTGCTGTGTGGCTCTCTTTGGGGAAG	3120
Db	3061	CTGCGAGCTATTATGACTTTTACAGCAAAACAAATTTGCTGTGTGGCTCTCTTTGGGGAAG	3120
Qy	3121	GGAAACAGGATAGCAGGAGGCTCAGGCTAGCAAGTCTGGAAGCTCAACCTTAAAGCCAGAGGCA	3180
Db	3121	GGAAACAGGATAGCAGGAGGCTCAGGCTAGCAAGTCTGGAAGCTCAACCTTAAAGCCAGAGGCA	3180
Qy	3181	TGGTTGATAGCAGAAAGTGAGGCTCTTCAACAGTGGGTGCTTAAGTAATCAGAAAC	3240
Db	3181	TGGTTGATAGCAGAAAGTGAGGCTCTTCAACAGTGGGTGCTTAAGTAATCAGAAAC	3240
Qy	3241	AGGAAGGCTCTGGTTGATGGAATTTATCAGTAAGATATCTACCCCTATCTCTCTCTCTAT	3300
Db	3241	AGGAAGGCTCTGGTTGATGGAATTTATCAGTAAGATATCTACCCCTATCTCTCTCTCTAT	3300
Qy	3301	AGAAAGCTAAACCGTCTCTCTCTCTTGTGTGTAGGCTGATATAACACGCTTGTCTTTTGTG	3360
Db	3301	AGAAAGCTAAACCGTCTCTCTCTCTTGTGTGTAGGCTGATATAACACGCTTGTCTTTTGTG	3360
Qy	3361	AGTGTTCAATGCTTTGCAAGATTTTTCAGTGTCTGCCAGTCTTCTGTAGAGGGTTGTGTAC	3420
Db	3361	AGTGTTCAATGCTTTGCAAGATTTTTCAGTGTCTGCCAGTCTTCTGTAGAGGGTTGTGTAC	3420
Qy	3421	CTTGACACCTGGGCTTGGATGTAGATGTCGCAAGGACACACTTCTGAAATGCCCTGTGTA	3480
Db	3421	CTTGACACCTGGGCTTGGATGTAGATGTCGCAAGGACACACTTCTGAAATGCCCTGTGTA	3480
Qy	3481	AAAGGTTATTATTCAATTTTACTTTGTCTTTGGAAAGGTGAAGTGTGTGGAAGAACTC	3540
Db	3481	AAAGGTTATTATTCAATTTTACTTTGTCTTTGGAAAGGTGAAGTGTGTGGAAGAACTC	3540
Qy	3541	ACAGAGATGATTCTCTGTAGGAAACCTTTTTTCCCTTAAAGCCTATATCCACT	3600
Db	3541	ACAGAGATGATTCTCTGTAGGAAACCTTTTTTCCCTTAAAGCCTATATCCACT	3600
Qy	3601	TTCAAGTCAACTTTTACATCTGTCATGAAAGAGTGTGTAGGCCGCTCT	3660
Db	3601	TTCAAGTCAACTTTTACATCTGTCATGAAAGAGTGTGTAGGCCGCTCT	3660
Qy	3661	CGTGGCTCTGGGAAAGCACCATAAGGGGAAGAAATGTTATGCCGAGAAATCTGACTGGC	3720
Db	3661	CGTGGCTCTGGGAAAGCACCATAAGGGGAAGAAATGTTATGCCGAGAAATCTGACTGGC	3720

Db	3661	CGTGGCTCTGGGAAAGCACCATAAGGGGAAGAAATGTTATGCCGAGAAATCTGACTGGC	3720
Qy	3721	AGGGAACCTGGTCTCAGAGCTCCCAAGACCACTACACAGTGTTTAAGTAGGAACAGTCCGAG	3780
Db	3721	AGGGAACCTGGTCTCAGAGCTCCCAAGACCACTACACAGTGTTTAAGTAGGAACAGTCCGAG	3780
Qy	3781	GGTGGGTTTCATATAATAGAAATGGAAACAGAGGGAGGGAAGATAGCTACAAAGTTTCTATAG	3840
Db	3781	GGTGGGTTTCATATAATAGAAATGGAAACAGAGGGAGGGAAGATAGCTACAAAGTTTCTATAG	3840
Qy	3841	GGTCCCTAAGTCTTTAAGATACAAAATAGCTGGTCTTCAATAACAAAGAAAGTCTGGG	3900
Db	3841	GGTCCCTAAGTCTTTAAGATACAAAATAGCTGGTCTTCAATAACAAAGAAAGTCTGGG	3900
Qy	3901	AAGCAGCAAGCAATTTGAGAGGGAGATGGAAGGGGAAAAAACAATGTAGAGGATTTGAAAA	3960
Db	3901	AAGCAGCAAGCAATTTGAGAGGGAGATGGAAGGGGAAAAAACAATGTAGAGGATTTGAAAA	3960
Qy	3961	GCTACAAATCCTCCAGAGAGGATTTTTCTTGGAGGAATCTAGAACAAAGGGTGTGGATT	4020
Db	3961	GCTACAAATCCTCCAGAGAGGATTTTTCTTGGAGGAATCTAGAACAAAGGGTGTGGATT	4020
Qy	4021	AGGTGGATCCAGAGGACTTGTCTTGGCCATTTGAATCTGGGTTTTGTCTCTCCATTGA	4080
Db	4021	AGGTGGATCCAGAGGACTTGTCTTGGCCATTTGAATCTGGGTTTTGTCTCTCCATTGA	4080
Qy	4081	GGTTGAGAGGCTCACCCCTTTTTTACCTGGATAGGAGGAGAAAGAGGGGTGTTTTGAC	4140
Db	4081	GGTTGAGAGGCTCACCCCTTTTTTACCTGGATAGGAGGAGAAAGAGGGGTGTTTTGAC	4140
Qy	4141	TCCTTACCTGAGTTTACTAGTTTACGCAATGGAAACAGACACTCGGACCTCTCTCTTGAC	4200
Db	4141	TCCTTACCTGAGTTTACTAGTTTACGCAATGGAAACAGACACTCGGACCTCTCTCTTGAC	4200
Qy	4201	AAGAAAAAAGAAAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAA	4260
Db	4201	AAGAAAAAAGAAAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAA	4260
Qy	4261	GCACAGGAGCTGGGATGCTGGCCCATGCTTTAAATCCAGCATTTGGGAGGAGAGGC	4320
Db	4261	GCACAGGAGCTGGGATGCTGGCCCATGCTTTAAATCCAGCATTTGGGAGGAGAGGC	4320
Qy	4321	AGGTGACTTTCTAAATTTCAAGGCGAGCTGCTGTCTACAAAGTGTCTTCAGGACAGCCAGG	4380
Db	4321	AGGTGACTTTCTAAATTTCAAGGCGAGCTGCTGTCTACAAAGTGTCTTCAGGACAGCCAGG	4380
Qy	4381	GCTATACAGAAACCTCTGTCTCGGAAAAAAGAAAAAGAAAGAAAGAAAGAAAGAAAG	4440
Db	4381	GCTATACAGAAACCTCTGTCTCGGAAAAAAGAAAAAGAAAGAAAGAAAGAAAGAAAG	4440
Qy	4441	AGAAAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAG	4500
Db	4441	AGAAAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAGGAGAG	4500
Qy	4501	AGGAGAGGAG	4560
Db	4501	AGGAGAGGAG	4560
Qy	4561	AAG	4620
Db	4561	AAG	4620
Qy	4621	AGAAAGAAAAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCA	4680
Db	4621	AGAAAGAAAAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCAAGCA	4680
Qy	4681	GGTCTTTTGAAGCAAGGCTTTTGAATTTGAGCGCTCATCAATAGTTGATCATGGTCAGGTGG	4740
Db	4681	GGTCTTTTGAAGCAAGGCTTTTGAATTTGAGCGCTCATCAATAGTTGATCATGGTCAGGTGG	4740
Qy	4741	AGGGCTACCTGTCTAGGCGAGCCCTGTGGCTTACCACTTAACTCTCCAGGCTCTCAGTA	4800
Db	4741	AGGGCTACCTGTCTAGGCGAGCCCTGTGGCTTACCACTTAACTCTCCAGGCTCTCAGTA	4800





Qy	892	ACCCACTGGGACATACATCTCTACTTACAGGCTTTCTTCCATCTCCTTGTCAACCAGG	951
Db	2571	ACCCACTGGGACATACATCTCTACTTACAGGCTTTCTTCCATCTCCTTGTCAACCAGG	2630
Qy	952	CACCTTAGGGTTTTCTCTCTTTTCAGGCCAGCCTTGAGATAACACACAGAGCTCCGGCTC	1011
Db	2631	CACCTTAGGGTTTTCTCTCTTTTCAGGCCAGCCTTGAGATAACACACAGAGCTCCGGCTC	2690
Qy	1012	ATCGGGGAGAAATCGTTCCGAGGAGTCAGTGTAAAGTCCTCACTGTGATGAGCAGGGCTAG	1071
Db	2691	ATCGGGGAGAAATCGTTCCGAGGAGTCAGTGTAAAGTCCTCACTGTGATGAGCAGGGCTAG	2750
Qy	1072	CTGCGGAGCTGGTGGACCTCTGGGATAGTCAGCTGATGACCCCTGCTCTCTTGTGTC	1131
Db	2751	CTGCGGAGCTGGTGGACCTCTGGGATAGTCAGCTGATGACCCCTGCTCTCTTGTGTC	2810
Qy	1132	TACCTGACGGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTCCTCAACTTCACCCCTGGAA	1191
Db	2811	TACCTGACGGCTAAGGATCAGTGTCTACCTGATGAAGCAGGTCCTCAACTTCACCCCTGGAA	2870
Qy	1192	GACATTCGTCTCCCGAGTCAGACAGGTTCCGGCCCTACATGACGAGGAGTGGTCCCTTTC	1251
Db	2871	GACGTTCTGTCTCCCGAGTCAGACAGGTTCCAGCCCTACATGACGAGGAGTGGTACCTTTC	2930
Qy	1252	CTGACCAAACTCAGCAATCAGCTCAGCTCCCTGCTGTAAGTCCTGGCTACCTATGCT	1311
Db	2931	CTGACCAAACTCAGCAATCAGCTCAGCTCCCTGCTGTAAGTCCTGGCTACCTATGCT	2990
Qy	1312	CCTCTCTCTCTCTTCTATTTCCAGTAAGAACCCGAGGTCCTGCCCCTCTCTCTCTCACAC	1371
Db	2991	CCTCTCTCTCTCTTCTATTTCCAGTAAGAACCCGAGGTCCTGCCCCTCTCTCTCTCACAC	3050
Qy	1372	AGAGTGAGGAGGCTCAGCAACCAACCATCATAGGCACTTGAAATGAGTCAAAAGG	1431
Db	3051	AGAGTGAGGAGGCTCAGCAACCAACCATCATAGGCACTTGAAATGAGTCAAAAGG	3110
Qy	1432	CTTTGGCTTCAATTTGAGTAATACCTTGTGATTTGATTTAGTTAAGCTTTATTTGTTTATC	1491
Db	3111	CTTTGGCTTCAATTTGAGTAATACCTTGTGATTTGATTTAGTTAAGCTTTATTTGTTTATC	3170
Qy	1492	CATGGAAGAAATCAACTCAAAATCTGTAGGATGAGAAAGATGTTGGGAACGAAAAAGG	1551
Db	3171	CATGGAAGAAATCAACTCAAAATCTGTAGGATGAGAAAGATGTTGGGAACGAAAAAGG	3230
Qy	1552	CCTAGATAGAAACAGATCTGTGATGATACAGTACTTATGAGGGGGGGGGGCGAGGGGCG	1611
Db	3231	CCTAGATAGAAACAGATCTGTGATGATACAGTACTTATGAGGGGGGGGGGCGAGGGGCG	3286
Qy	1612	ATATCCACTGAGTCCAAAGTACTTGTGGGAGAGAAATCCACTGAGTACAGTACTTGT - -	1669
Db	3287	ATATCCACTGAGTACAAAGTACTTGTGGGAGAGAAATCCACTGAGTACAGTACTTGTGTG	3346
Qy	1670	-----GGGGGAAGGAATGGCACAGACAAAGT	1697
Db	3347	GCATGGAGATCCACTGAGTACAAAGTACTTGTGGGGGAGGGNAITGGCACAGACAAAGT	3406
Qy	1698	TGAAGGGAAGAGGAAGATGAGAGGCTCAATGTTGGGGGTGTGAAAGGTCACTCCTTT	1757
Db	3407	TGAAGGGA - - -AGGAAGATGAGAGGCTCATGTTGGGGGTGTGAAAGGTCACTCC - TT	3462
Qy	1758	TTCCATGTGATGGAGAGTTAGAAAAATCAGTGTGATGTTGATGTTCTTCAGACACCCC	1817
Db	3463	TTCCATGTGATGGAGAGTTAGAAAAATCAGTGTGATGTTGATGTTCTTCAGACACCCC	3522
Qy	1818	AA-----CTATGCGACACTGTGGGAGACCTGGCATTTAGGGAA	1855
Db	3523	CAACTATGAACATATCAACGAGGAGCGGCGAGACTGTGGGAGACCTGGCATTTAGGGAA	3582
Qy	1856	GGCGGGCTTTTCAACGAGAAACCTTATGCTCATCTCTGTGCTPACACTCCACCTTTG	1915
Db	3583	GGCGGGCTTTTCAACGAGAAACCTTATGCTCATCTCTGTGCTPACACTCCACCTTTG	3642

Qy	1916	ATGAGGTAAAGCTCAGGTTTCGTTTCTTACCGTCTTCTGCTACTGCTGGAACCTTCAGTAGG	1975
Db	3643	ATGAGGTTCAGCTCAGGTTTCGTTTCTTACCGTCTTCTGCTACTGCTGGAACCTTCAGTAGG	3702
Qy	1976	ATTCCCCAAAGACGAGGACAGCTCTTCTGTAAAGGAGGGAACCTGGGATTCAGTGTCTCTAG	2035
Db	3703	ATTCCCCAAAGACGAGGACAGCTCTTCTGTAAAGGAGGGAACCTGGGATTCAGTGTCTCTAG	3762
Qy	2036	AGAACGAATAGCTCAGAGGATCTTAGGTCAACGTGAAATCTTAGGTCAACGCGGCAAAAA	2095
Db	3763	AGAACGAATAGCTCAGAGGATCTTAGGTCAACGTGAAATCTTAGGTCAACGCGGCAAAAA	3822
Qy	2096	TGACTGAACCCCTCTATTTCCAGGTGAACCGCTCAGTGCCTCAGATATACCTAGGATATTGG	2155
Db	3823	TGACTGAACCCCTCTATTTCCAGGTGAACCGCTCAGTGCCTCAGATATACCTAGGATATTGG	3882
Qy	2156	GCTCCCAACCGGATAAGATTCGTGTAGTGTCTGTCTTTTATTTTTCGAGCACATCAGTGGT	2215
Db	3883	GCTCCCAACCGGATAAGATTCGTGTAGTGTCTGTCTTTTATTTTTCGAGCACATCAGGCGT	3942
Qy	2216	GACGACGAGAACATCCAGAGAGATGTCAGAGGCTGAAGGACAGTGAAAGAGTACTA	2275
Db	3943	GACGACGAGAACATCCAGAGAGATGTCAGAGGCTGAAGGACAGTGAAAGAGTACTA	4002
Qy	2276	TTGGCAAGCCCAATATTAAGCCATTCAGTAGGAGACGTGGGGAATTTCTTCTCTGCTTC	2335
Db	4003	TTGGCAAGCCCAATATTCAGTCCATTCAGTAGGAGACGTGGGGAATTTCTTCTCTGCTTC	4062
Qy	2336	CCAGTCTCTCTACTTTTGTAAACATTTTCTTTTGAACCTGTCTACTCTGTCTGCTGCTCA	2395
Db	4063	CCAGTCTCTCTACTTTTGTAAACATTTTCTTTTGAACCTGTCTACTCTGTCTGCTGCTCA	4122
Qy	2396	CTTAGCTGCACTGCATCTAGCTGGGCTATAGATCTTTCAATCTGTCTCTTAAATTTGTA	2455
Db	4123	CTTAGCTGCACTGCATCTAGCTGGGCTATAGATCTTTCAATCTGTCTCTTAAATTTGTA	4182
Qy	2456	AGTCACAATTCAGGAGCTAGCAGAAAGCTTAGCTCAGCAGCTCTCATGAGCACCTTGCTCG	2515
Db	4183	AGTCACAATTCAGGAGCTAGCAGAAAGCTTAGCTCAGCAGCTCTCATGAGCACCTTGCTCG	4242
Qy	2516	GAGGATGGCTGTGACAGAGTCAATGTAGAAAGACAGATCCCTGATTTCCAGCTCTGCA	2575
Db	4243	GAGGATGGCTGTGACAGAGTCAATGTAGAAAGACAGATCCCTGATTTCCAGCTCTGCA	4302
Qy	2576	CTTCCCTAGTGGCCACCTGTAAATTTACTTTAGCCCTGATTAGTATTTTGGAAAGCCAAATTC	2635
Db	4303	CTTCCCTAGTGGCCACCTGTAAATTTACTTTAGCCCTGATTAGTATTTTGGAAAGCCAGTTTC	4362
Qy	2636	CCACCGACCTACATAATCCGAAAGACGATGCATTTGAAAACTAGAAAAAGCTGGGCAAACT	2695
Db	4363	CCACCGACCTACATAATCTGNAGAACCAATGCATTTGAAAACTAGAAAAAGCTGGGCAAACT	4422
Qy	2696	TACTAGAGATGATTTTGGAGCTCAATAAACCTGATGCTCTGAAATGTGATCAAAATCAACCC	2755
Db	4423	TACTAGAGATGATTTTGGAGCTCAATAAACCTGATGCTCTGAAATGTGATCAAAATCAACCC	4482
Qy	2756	AGAAATACAAAGAGCTGGATTTGCAATAGAGCAAGTATTTAGAAATCACTGGGTATT	2815
Db	4483	AGAAATACAAAGAGCTGGATTTGCAATAGAGCAAGTATTTAGAAATCACTGGGTATT	4542
Qy	2816	AAAGCTGTCTATTTAAATATAATAGTGTCTATTTAGCTGCCCTATTAAAGATTTAAACA	2875
Db	4543	AAAGCTGTCTATTTAAATATAATAGTGTCTATTTAGCTGCCCTATTAAAGATTTAAACA	4598
Qy	2876	CAAGAGTGGATTAATCTTCCCAATTTACTGGGCTCGTTTCAATAGAGTAAATATACAGTC	2935
Db	4599	CAAGAGTGGATTAATCTTCCCAATTTACTGGGCTCGTTTCAATAGAGTAAATATACAGTC	4658
Qy	2936	ATAGATTAATTTATAGTGTCAAGAGTATGATGTTGGAAACCTTTCTCTTACTTTTTACT	2995
Db	4659	ATAGATTAATTTATAGTGTCAAGAGTATGATGTTGGAAACCTTTCTCTTACTTTTTACT	4718
Qy	2996	TCAATTTCTTAGTTATTTATTTTCTTTCACACCTGATCAAGCCACTAGTAGACACT	3055



QY 5212 CTTATTTTCATAGCTTCGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCT 5271  
Db |||||||  
QY 6526 CTTATTTTCATAGCTTCGAGAGAGTGAGAGATCAAGCGGATGGGAACTGGACCTGCT 6585  
Db |||||||  
QY 5272 GTTTATGCTCTGAGAAATGCTTCGGTCTGAGCGAGAGAGCTAGAAAACGAAAGACTG 5331  
Db GTTTATGCTCTGAGAAATGCTTCGGTCTGAGCGAGAGAGCTAGAAAACGAGAACTG 6645  
QY 5332 CTCCTTCCTGCTTTCTAAAAGAAACAATAGATCCCTGAATGGACTTTTAAAGGA 5391  
Db |||||||  
QY 6646 CTCCTTCCTGCTTTCTAAAAGAAACAATAGATCCCTGAATGGACTTTTAAAGGA 6705  
Db |||||||  
QY 5392 AAGTGAGAGCTACGTCACCATCAATAGAGATTTCACATGAAACCTGCTCAGTTGA 5451  
Db AAGTGAGAGCTACGTCACCATCAATAGAGATTTCACATGAAACCTGCTCAGTTGA 6765  
QY 5452 AAGAGAAATAGTGTCAAGTTGTCCATGAGACCAAGAGGTAGACTTGTATAACCAAGAT 5511  
Db AAGAGAAATAGTGTCAAGTTGTCCATGAGACCAAGAGGTAGACTTGTATAACCAAGAT 6825  
QY 5512 TCATGTCAATATTTTATGTGATGATAATGCAACAGAAAAAGATGATGACTTTAAAAA 5571  
Db TCATGTCAATATTTTATGTGATGATAATGCAACAGAAAAAGATGATGACTTTAAAAA 6885  
QY 5572 TTGTTTGAAGAGGTACCTCTCATTCCTCTAGAGAAAGCCCTATGTAACCTTCATTC 5631  
Db TTGTTTGAAGAGGTACCTCTCATTCCTCTAGAGAAAGCCCTATGTAACCTTCATTC 6945  
QY 5632 CATAACCAATCTTTATATATGTAAGTTTATTTATATAGTATACATTTTATTTATGTC 5691  
Db CATATCCNAATTTTATATATGTAAGTTTATTTATATAGTATACATTTTATTTATGTC 7005  
QY 5692 AGTTTATTAATGGAATTTATTTATAGAAAAATTTATCTGATGTTGATTTGATATAA 5751  
Db AGTTTATTAATGGAATTTATTTATAGAAAAATTTATCTGCTATTGATATT-AGTATAAG 7064  
QY 5752 GCAATAATATTTATGATAATTAACATAGAAACAGATATCTTAGGCTTTTAAATAACACA 5811  
Db GCAATAATATTTATGACAAATTAACATAGAAACAGATATCTTAGGCTTTTAAATAACACA 7124  
QY 5812 TGAATATCATAAATCTCTGCTCTGTGTAATTTTCTCCCTTAAATATCAACAATACCATCA 5871  
Db TGGATATCATAAATCTCTGCTCTGTGTAATTTTCTCCCTTAAATATCAACAATACCATCA 7184  
QY 5872 TCGTCATCATTTACCAATCATTTCTCATGACTTTCATGCTGTGATCTCATATATCTGGTAAAG 5931  
Db TCATCATCATTTACCAATCATTTCTCATGATTTTCATGCTTTGACCCATATTAATCTGTTAAA 7244  
QY 5932 TTTG 5935  
Db |||||  
7245 GTTG 7248

RESULT 4  
US-10-627-273-8  
; Sequence 8, Application US/10627273  
; Publication No. US2004010189A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/10/627,273  
; CURRENT FILING DATE: 2003-07-25  
; PRIOR APPLICATION NUMBER: US/09/751,797  
; PRIOR FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 8  
; LENGTH: 7445  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-10-627-273-8

Query Match 71.5%; Score 4245.2; DB 7; Length 7445;  
Best Local Similarity 88.3%; Pred. No. 0;  
Matches 5039; Conservative 0; Mismatches 178; Indels 487; Gaps 20;

QY 293 TACCATCTATCCGACGAGCATGTTCCCTGTGATGTTTTTGCCTTTTGCCTCTCTCGCTAAC 352  
Db |||||||  
QY 1971 TACCATGCTACCCGACGAAACATGCTCCCTGATGTTTTTGCCTTTTGCCTCTCTCACATAAC 2030  
Db |||||||  
QY 353 AGGCTCTCTCTCAGTTATCAACTTTTGAACATTTGCGGATCGGTGATGCGTGTCTTGC 412  
Db AGGCTCTCTCTCAGTTATCAACTTTTGAACATTTGCGGATCTCTGATGGCTGTCTCTGCA 2090  
QY 413 GAAATCTATGAGTTTTTCCCTTATGGGACTTTTGGCCGCGAGCTGCTTCTTCATTGTC 472  
Db GAAATCTATGAGTTTTTCCCTTATGGGACTTTTGGCCGCGAGCTGCTTCTTCATTGTC 2150  
QY 473 CCTGTGGGCGCAGGAGCAAAATGCGTGCCTCATCAACACCGGCTGCAAGCTTGAAGTGT 532  
Db CCTGTGGGCGCAGGAGCAAAATGCGTGCCTCATCAACACCGGCTGCAAGCTTGAAGTGT 2210  
QY 533 CAATCTCCAGCGCTGATCATGCTCAACCGGCACTTTATGCTGGCCAAAGAGGTACAGCT 592  
Db CAATCTCCAGCGCTGATCATGCTCAACCGGCACTTTATGCTGGCCAAAGAGGTACAGCT 2270  
QY 593 GCATCTCTTCTCTCCATACGGCTTGCCA-TTCTCTGAGCACTTTGCAAACTCTTTAG 651  
Db GCATCTCTTCTCTCCATACGGCTTGCCA-TTCTCTGAGCACTTTGCAAACTCTTTAG 2330  
QY 652 GGGCGCTTTATCTCCGAGGCTCTCACTACCTATGTTTCTGTCTCTTTAGAGACTCTTTA 711  
Db GGGCGCTTTATCTCCGAGGCTCTCACTACCTATGTTTCTGTCTCTTTAGAGACTCTTTA 2390  
QY 712 AGGACTGGATCTTTTCTATTTTCAAGGCTCTCAGGACCAATTTCTTATCTTGGCT 771  
Db AGGACTGGATCTTTTCTATTTTCAAGGCTCTCAGGACCAATTTCTTATCTTGGCT 2450  
QY 772 TCAGGACACATATCTGAATTTTATCTACAGAGCGGCTTTAGAAAGCCACCCAGACTG 831  
Db TCAGGACACATATCTGAATTTTATCTACAGAGCGGCTTTAGAAAGCCACCCAGACTG 2510  
QY 832 CAATACCTTTCCATCTGTGTGCTCTCTCTGAACTCATCTCTTTGGCTACTCTTGAG 891  
Db CAATACCTTTCCATCTGTGTGCTCTCTCTGAACTCATCTCTTTGGCTACTCTTGAG 2570  
QY 892 ACCACCTGCGGACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTTGTACCCAGG 951  
Db ACCACCTGCGGACATACATCTCTACTTACAGGCTTTTCTTCCATCTCTTGTACCCAGG 2630  
QY 952 CACTTAGGGTTTTCTCTTTTCAGGCCAGCTTCAGATACACACAGAGCTCGGGCTC 1011  
Db CACTTAGGGTTTTCTCTTTTCAGGCCAGCTTCAGATACACACAGAGCTCGGGCTC 2690  
QY 1012 ATCGGGGAGAAATGTTTCCGAGGAGTCAGTGTAAGTCTCTCACTGTGATGAGCAGGGCTAG 1071  
Db ATCGGGGAGAAATGTTTCCGAGGAGTCAGTGTAAGTCTCTCACTGTGATGAGCAGGGCTAG 2750  
QY 1072 CTGCGGAGCTGTGTGAGCCCTCTCGGATPAGTCTGACGTATGACCCCTGCTGCTTCTGTC 1131  
Db CTGCGGAGCTGTGTGAGCCCTCTCGGATPAGTCTGACGTATGACCCCTGCTGCTTCTGTC 2810  
QY 1132 TACCTGCAGGCTAAGGATCAGTGTCTCTGATGAGCAGGCTCAACTTCCACCTCGAA 1191  
Db TACCTGCAGGCTAAGGATCAGTGTCTCTGATGAGCAGGCTCAACTTCCACCTCGAA 2870  
QY 1192 GACATTTCTGCTCCCCCAGCTCAGACAGGTTCCGGGCCCTACATGACAGGAGGTGGTGCCTTTC 1251



QY	3356	TTTTGAGTGTTCATGGCTTTGCAGATTTTTCAGTGTCTGCCAGTTCCTGTTAGAGGGTTT	3415
Db	5070	TTTTGAGTGTTCATGGCTTTGTAGATTTTGTAGTGTCTGCCAGTTCCTGTTAGAGGGTTT	5129
QY	3416	GTTTACCTTGACACCTGGGCTTGGATGTTAGCATGCCAAAGGCAACACTTCTGAATGCCCT	3475
Db	5130	GTTTACCTTGACACCTGGGCTTGGATGTTAGCATGCCAAAGGCAACACTTCTGAATGCCCT	5189
QY	3476	GTGTAAAGGTTATTATTCAATTTACTTTGTCTTTGGAAAGGTGAAGTGTGTGAGAAAG	3535
Db	5190	GTGTAAAGGTTATTATTCAATTTACTTTGTCTTTGGAAAGGTGAAGTGTGTGAGAAAG	5249
QY	3536	AACTTCACAGGAGATGTAATCTCTGTAGGAAAC- TTTTTTTTTCCCTTTAAAGCCCTATAA	3594
Db	5250	AACTTCACAGGAGATGTAATCTCTGTAGGAAAC- TTTTTTTTTCCCTTTAAAGCCCTATAA	5309
QY	3595	TCCACTTTCAGTCAACTTTGACTTTTATACATGCTGTCTCATGAAAGAGTGTTAGGCC	3654
Db	5310	TCCACTTTCAGTCAACTTTGACTTTTATACATGCTGTCTCATGAAAGAGTGTTAGGCC	5369
QY	3655	CGCTCTCGTGGCTCTGGGAAAGCACCAATAGGGGAGAAATGTTATGCCGAGAAATCTG	3714
Db	5370	CGCTCTCATGGCTCTGGGAAAGCACCAATAGGGGAGAAATGTTATGCTGAGAAATCTG	5429
QY	3715	ACTGGCAGGGAACCTGGGTCAGAGCTCCCCAAAGACCACCTACAGGTGTTTAAGTAGGAACA	3774
Db	5430	ACCGGCAGGGAACCTGGTCAGAGCTCCCCGAGACCCACACAGGTGTTTAAGTAGGNACA	5489
QY	3775	GTCAGGGTGGGTTTCATATATAGAAATGGAACAGAGGGAGGAAGATAAGCTCAAAAGTT	3834
Db	5490	GTCAGGGTGGGTCATGTAATAGAAATGGAACAGAGGGAGGAAGATAAGCTCAAAAGTT	5549
QY	3835	TCATAGGTCCTAAGTCTTTAAGNATACAAATAGCTGGTGGGCTTCATACCAAGGAAG	3894
Db	5550	TCATAGGTCGGGAGTCTTAAAGNATACAAATAGCTGGTGGGCTTCATACCAAGGAAG	5609
QY	3895	TCTCGGAAGGCAGCAAGCATTTGAGGGGAGATGGAAGGCAAAAAAAC- AATGTAGAGGA	3952
Db	5610	TCTCGGAAGGCAGCAAG- -TGAGGGGAATGGAAGGGAAGAAAAACAGAAATGTAGAGGA	5666
QY	3953	TTTGAAAGCTACAAATCCTCCACGAGAGGATTTTCTCGAGGAATCTAGAACAAAGGTT	4012
Db	5667	CTTGAAACAGCTACAAATCCTCTACCAGACGATTTTCTTGGAAACAATCTAGAA- -GGT	5722
QY	4013	GGTGGATTAGGTGGATCGCAGAGGACTTGTCTTGCCATTGTAATCTGGGTTTTTGTCTC	4072
Db	5723	AGTGGATTAGGT- GAITTCAGGGGGACTTGTCTTGCCATTGTAATCTGGGTTTTTGTCTC	5781
QY	4073	TCCATTGAGGTTGAGAGCGTCACCCCTTTTACCCTCGATAGGAGGAGGAAGAGGGGT	4132
Db	5782	TCCATTGAGGTTGAAAGCGTCACCC- TTTTACCCTCGAATGGAGGAGGAAGAGGGGT	5840
QY	4133	GTTTTGACTCTACTCGGAGTTTTTACTAGTTTACGCAATGGAACAGACACTCGGACCTC	4192
Db	5841	GTTATGACTCTACTCGGAGTTTTTACTAGTTTACGCAATGGAACAGACACTCGGACCTC	5900
QY	4193	CTCTTGACAGAAAAAAGAAAAAGAAACCTGTGTTTCTCTTGTGTTTCTTTT	4252
Db	5901	CTCTTGAC- -AAAAAAGTGAACCTGTGTTTGTGTTTCTTTTGTGTTT	5950
QY	4253	TTAAGAGGACGAGCGAGCTGGGCATGGTGGCCCATGCCCTTAATCCAGCATTTGGGAG	4312
Db	5951	TTAAGAGGACAA- -	5963
QY	4313	GCAGAGGAGGTGACTTTCTTAAATTTCAAGGCCAGCCTGGTCTACAAAGTGGTTCCAGGA	4372
Db	5964	- - - - -	5963
QY	4373	CAGCCAGGCTATACAGAGAAACCTGTCTCGGAAAAAAGAAAAAGAAAGAAAG	4432
Db	5964	- - - - -	5963







Db 1282 AGGTTGCGTAA---GATGAGAAAGGTGTGGGAAAACATCTAGCTGTGGAAAATGGAATCCA 1338  
 Qy 1619 CTGAGTCCAAAGTACTTGTGGGAGAGAAATCCACTGAGTACAGTACTTGTGGGGGAAGG 1678  
 Db 1339 TTGAGTCTAAAGTGTGGGGGAGGGATGGCATGGAGAGAAATTAGAAGAGAAAGTGGG 1398  
 Qy 1679 AATGGCACAGACGAAAGTTCAGGGAAGAGGAGAGATGAGAGGCGCTCAATGTTGGGG 1738  
 Db 1399 AATGGGAAGGCTTAAAGTGC-----GTGGTGGGTGGGACACTGTTGCCCTGTTGA 1450  
 Qy 1739 TGTGAAAAGGTCACCTCTTTTCCATGTGATGGAGAGTAAAGAAAATCAGTGTGTGAGTT 1798  
 Db 1451 TGTGATGGGAGCCACAAAATCGGAGCGGTGTGAATCTGATGCCGCTGAACAATTTGAAC 1510  
 Qy 1799 TGATGCTTTCAGACACCCCAACTATGAGGAGACTGTGGGAGACCTGGGCAATTTAGGGA-AGG 1857  
 Db 1511 TATGAAAAAAGTGTGAGTGGAGTGGGCCCAAGTAAAAGGCCCTAGGACCTTACTGAAGAGG 1570  
 Qy 1858 CGCGGCTTTTCACAGGAGAACTTTATGCTCATCTCTGTGCTACACTCCACACCTTTGAT 1917  
 Db 1571 GCTTAAATTTTCACATGAGATGTTTTATGTACATTTCTTTGTTCTAAGCATGCAATTTCTG 1630  
 Qy 1918 GAGGTAAAGCTCAGGTTTCGTTTCT-----ACCGTTCTTGTGCTAC 1956  
 Db 1631 GAGATACGATGAGGTTTATTCCTACAGAAATTTGCATAAACTACTCCGCTCTTTCCAC 1690  
 Qy 1957 TGGTGGAAACTTTCAGTAGGATTCCTCCAAAGACGAGGACAGCTCTTCTGTAAAGGAGGAC 2016  
 Db 1691 AAATGCAAACTCAGTAGGATTTCCAAAGATGAAGAGAGGCTCTTTGTAAGGGAAGTGA 1750  
 Qy 2017 CTGATTTTCAGTGTCTTAGAGAACAAATAGCTCAGAGAACTAGGTCAACGTCGAAATCT 2076  
 Db 1751 CTGATTTCTGCGTCCAGGGAATTCAGAGCTCAGGAAATCTAGGTCACCTGTTGGAATC 1810  
 Qy 2077 AGGTACAGCGGGCAAAATGACTGAACGCTCTATTCCAGGTGAACGGTCACGTGCGTC 2136  
 Db 1811 TAGGTCATTTGGGCAAAATTTACTAAGAGCTTTAATTCAGGTGAATGTACTGTACCTC 1870  
 Qy 2137 AGATATAGTAGGATTTGGGGTCCCAACGGATATAGATTTCTGTTAGTGA-GTCTGCTTTTA 2195  
 Db 1871 CATGGGTGGAGGTTTCATAAAGTTTCAGCAACAATTAAGATAGTTATGCTGTTATTG 1930  
 Qy 2196 TTTTTCAGCACATCAGTGTGACAGCAGACATCCAGAAATGTCAGAGGCTGAAGG 2255  
 Db 1931 TTTTATAGCATTTGAAGGTGATGACCTGTCATATCCAGAGGAATGTGCAAAAGCTGAAGG 1990  
 Qy 2256 AGACAGTGAAGAAAGTACTATTGGCAAGCCACAATCTAAGCCATTCAGTAGGAGACGTG 2315  
 Db 1991 ACACAGTGAAGAAAGTAGGACTGATATCTGTCATGCTAAGTCATGCAATAGGAGAGACA 2050  
 Qy 2316 GGGATTTCTTCTGCTTCCAGTCTCTTCT--ACTTTGTAACAATTTTCTTGTGACTTGT 2373  
 Db 2051 AATGTTGTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT 2110  
 Qy 2374 CTACTGCTGCTCCATTTACTCATTAGCTGACCTGCACTGAGCTGAGCTGGGCTATAGATCTT 2433  
 Db 2111 CTACACAGGGCGAAT----ACTTTGGTGTCTGTATGTAGATATATCTATATATCTA 2166  
 Qy 2434 TCAATCTGTCTTAAATTT---GTAAGTCACAATTTCTGGAGCTAGCAGAAAGCTTAGCTC 2490  
 Db 2167 GATGTCAGTTCCAAATCTTCGAAATTTAGNAATTTCTAGAACTGGTTGGGATCTTAGCTT 2226  
 Qy 2491 AGCCAGTCTCATAGACATCTGCTGGAGGATGGCTTTGTGACAGAGTCAATGCTTAGAGAC 2550  
 Db 2227 GTCTAGTCACATAACCTCAGATTTCTGGGGATGGTCAGTGGCAGAGATAGGGCTAGAATGC 2286  
 Qy 2551 AGCATCCCTGATTTCCAGCTCTGAC- TTGCTAGTGGCCAGCTGTAATTTACTTTAGCCT 2609  
 Db 2287 AGGTCTCTGAATCCCAAGCAGCACTTTTCCCGTGGTGATACAGATTAGTTTGGTAC 2346  
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Db 2347 CATTAATTTCTTAGGAAAATTTCCAGATTTCTTATTTGACTCATGTATCTGAAGAAGTACTTG 2406  
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 Db 2407 TTTAAAAACAGAAAATATGCTATGGGCAATTTATTTTGAAGTCATTTTTGAAGTCAATA 2466  
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 Db 2467 TGCATTTGCTTTGAAACTTGGGAAGATAAATCTCAGAACAAATGAGAAAAGAGCTGGACTGC 2526  
 Qy 2784 AANTAGACAAGTATTTAGAAATCAGCTGGTATTTAAACAGCTGTCTATTTAATTAATAATAG 2843  
 Db 2527 ATATAGGCTTAAATTTCTGGAGTAAATAAACACTTAT-----TTTGAATTTATCATAATA 2578  
 Qy 2844 TGTCTATTTAGCTGCCCTATTTAAGATTTAAACACAGAGTGGATAACTTCCCAATTTACTG 2903  
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 Qy 2904 GGCTCTGTTTCAATAGAGTAAAAATATCAGTCTATAGATTAATATATAGTGTATGAAAGTA 2963  
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 Qy 2964 TGAGTTGGAACCC---CTTTCCCTTACTTTTACCTTCACTTCTTAGTATTTATTTTCTT 3020  
 Db 2695 TGAATTTGTTAAGCTTTTTTTTTTCTTCTCTCTCCCATCAAGACCTTCCATTTAGTTCTT 2754  
 Qy 3021 TCTTCAACCCCTGATCAAGCCACTAGTAAGCACCTATCTGTCGAGCTATTAATATGACT 3080  
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 Db 2815 CTATAGTGAATGATACCATCATGTGGCTTATTTGGTGAAAGAAACA--ACAATGGAAGGC 2872  
 Qy 3141 TCAGGCTAGCAGTCTGAGACTCAACCTAAAGCCAGAGCATGTTGTATAGCAGAGAAAGT 3200  
 Db 2873 TTAGACTTAACAATAGT-GACTCACCCCAAAACCGAGGAAATGATTAGGAGCAGTGAAGT 2931  
 Qy 3201 GAGGCTCTTCAACAGTGGGTGTGCTTAAGTAATCAGAAAAACAGGAAGGCTCTGGTTGATGG 3260  
 Db 2932 GAGGCTCTT-GCAAGCAGGTACAACTAAATCTCAGAAACATGAAGGCTCAGTTGATGG 2990  
 Qy 3261 AATTATCAGTAAGATATCTACCCCTTATCTCC-----TTCTCTATAGAGCTTAAACCG 3313  
 Db 2991 AATTTTCAGTAACAGCTTAACCTTAATTTCCCTCTTCTGACTTTTAAAAAA 3050  
 Qy 3314 TCTCTCTCTCTGTGTGTAGGCTGATAAACACGCTTGT--TTCTTTTGTAGTGTTCATGG 3371  
 Db 3051 GCGTTTCTTCTGAGCATCATTTAATGAGTGTGACTGTTTCTCTTCTTGAATAATGGAAG 3110  
 Qy 3372 CTTTGCAGATTTTTCAGTGTCTGCGAGTCTTGT--TAGAGGGTTTGTACTTTGACACC 3429  
 Db 3111 CTTTGTAGTTTAAATTTGTGAGCCAGTCTCTTGTATAGAACTATTTATAGACATG 3170  
 Qy 3430 TGGCTTGGATTTAGCATGCCAAAGGCACACACTTCTGAATGCTGTGTAAGAGTTAT 3489  
 Db 3171 GAGGCTGAAATTTAGCATGCCACAGACAGGCGATGCTTTACATCTTGTCTTAAAAAAT 3230  
 Qy 3490 TATTCATTTACT-----TTGTCTTTGGAAGGTGAAGTGTGTGAGAAAGACTCA 3541  
 Db 3231 TACTGATTTCACTTGTCTTGTGTTGTTCTTTAGAAAAAGTGAAGTGTGAGAGAGGAATCTCA 3290

RESULT 6

US-10-627-273-25  
 ; Sequence 25, Application US/10627273  
 ; Publication No. US20040110189A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Dumoutier, Laure  
 ; APPLICANT: Renauld, Jean-Christophe  
 ; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
 ; TITLE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof





Db 808 ATATTTTATTTGTCATTGATATGCAACAGAAAAAGTATGACTTTTAAAAAATTTGTTGAA 867  
QY 5581 AGGAGGTTTACCTCTCATTTCTCTAGAGAAAAGCCTATGTAACCTTTCAATTTCCATAACCAA 5640  
Db 868 AGGAGGTTTACCTCTCATTTCTCTAGAGAAAAGCCTATGTAACCTTTCAATTTCCATAACCAA 927  
QY 5641 TACTTTATATATGTAAGTTTATTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700  
Db 928 TACTTTATATGTAAGTTTATTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 987  
QY 5701 ATATGGAATTTTATTAAGAAAATTTATCTGATGTTGATATTTGAGTATAAGCAATAAT 5760  
Db 988 ATATGGAATTTTATTAAGAAAATTTATCTGATGTTGATATTTGAGTATAAGCAATAAT 1047  
QY 5761 ATTTATGATAATACTATAGAAAACAAGATATCTTAGGCTTTTAAATAACACATGAATATCA 5820  
Db 1048 ATTTATGATAATACTATAGAAAACAAGATATCTTAGGCTTTTAAATAACACATGAATATCA 1107  
QY 5821 TAAA 5824  
Db 1108 TAAA 1111  
RESULT 8  
US-10-627-273-9  
; Sequence 9, Application US/10627273  
; Publication No. US20040110189A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFS) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/10/627,273  
; CURRENT FILING DATE: 2003-07-25  
; PRIOR FILING DATE: US/09/751,797  
; PRIOR FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-10-627-273-9  
Query Match 10.1%; Score 602.4; DB 7; Length 1111;  
Best Local Similarity 99.8%; Pred. No. 2.7e-123;  
Matches 603; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
QY 5221 ATAGCTTGGAGAGCGGAGAGATCAAAGCGATCGGGGAACCTGACCTGCTGTTTATGTC 5280  
Db 508 AAAGCTTGGAGAGCGGAGAGATCAAAGCGATCGGGGAACCTGACCTGCTGTTTATGTC 567  
QY 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGAACTGCTCCTTCCT 5340  
Db 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGAACTGCTCCTTCCT 627  
QY 5341 GCCTTCTAAAAGAACAAATAAGATCCCTGAATGACCTTTTACTTAAAGGAAAGTGAAG 5400  
Db 628 GCCTTCTAAAAGAACAAATAAGATCCCTGAATGACCTTTTACTTAAAGGAAAGTGAAG 687  
QY 5401 GCTAACGCTCCACCATCATTTAGAGATTTCATGAAACCTGGCTCAGTTGAAAGAGAAA 5460  
Db 688 GCTAACGCTCCACCATCATTTAGAGATTTCATGAAACCTGGCTCAGTTGAAAGAGAAA 747  
QY 5461 TAGTGTCAGTTGTCATGAGACGAGAGGTAGACTTTGATTAACCAACAAAGATTCATTGACA 5520

Db 748 TAGTGTCAGTTGTCATGAGACGAGAGGTAGACTTTGATTAACCAACAAAGATTCATTGACA 807  
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Db 808 ATATTTTATTTGTCATTGATATAATCAACAGAAAAAGTATGACTTTTAAAAAATTTGTTGAA 867  
QY 5581 AGGAGGTTTACCTCTCATTTCTCTAGAGAAAAGCCTATGTAACCTTTCAATTTCCATAACCAA 5640  
Db 868 AGGAGGTTTACCTCTCATTTCTCTAGAGAAAAGCCTATGTAACCTTTCAATTTCCATAACCAA 927  
QY 5641 TACTTTATATATGTAAGTTTATTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700  
Db 928 TACTTTATATATGTAAGTTTATTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 987  
QY 5701 ATATGGAATTTTATTAAGAAAATTTATCTGATGTTGATATTTGAGTATAAGCAATAAT 5760  
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QY 5821 TAAA 5824  
Db 1108 TAAA 1111  
RESULT 9  
US-09-751-797-7  
; Sequence 7, Application US/09751797  
; Patent No. US20010024652A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Louhed, Jamila  
; APPLICANT: Renauld, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: (TIFS) The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LUD 5543.2  
; CURRENT APPLICATION NUMBER: US/09/751,797  
; CURRENT FILING DATE: 2000-12-29  
; PRIOR APPLICATION NUMBER: 09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 29  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-09-751-797-7  
Query Match 9.4%; Score 555.2; DB 3; Length 1119;  
Best Local Similarity 96.0%; Pred. No. 8.9e-113;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
QY 5221 ATAGCTTGGAGAGCGGAGAGATCAAAGCGATCGGGGAACCTGACCTGCTGTTTATGTC 5280  
Db 510 AAAGCTTGGAGAGCGGAGAGATCAAAGCGATCGGGGAACCTGACCTGCTGTTTATGTC 569  
QY 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGAACTGCTCCTTCCT 5340  
Db 570 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAAAGAACTGCTCCTTCCT 629  
QY 5341 GCCTTCTAAAAGAACAAATAAGATCCCTGAATGACCTTTTACTTAAAGGAAAGTGAAG 5400  
Db 630 GCCTTCTAAAAGAACAAATAAGATCCCTGAATGACCTTTTACTTAAAGGAAAGTGAAG 689  
QY 5401 GCTAACGCTCCACCATCATTTAGAGATTTCATGAAACCTGGCTCAGTTGAAAGAGAAA 5460  
Db 690 GCTAACGCTCCACCATCATTTAGAGATTTCATGAAACCTGGCTCAGTTGAAAGAGAAA 749  
QY 5461 TAGTGTCAGTTGTCATGAGACGAGAGGTAGACTTTGATTAACCAACAAAGATTCATTGACA 5520

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Db 750 TAGTGTCAGGTTGTCATGAGACGAGGTTAGACTTGTGATTAACCAAGAGATTCATTGACA 809
Qy 5521 ATATTTTATTTGTCATGATTAATGCAACAGAAAAGTATGACTTTTAAAAAATTGTTTGA 5580
Db 810 ATATTTTATTTGTCATGATTAATGCAACAGAAAAGTATGACTTTTAAAAAATTGTTTGA 869
Qy 5581 AGGAGGTTTACCTCTCATTCTCTAGAGAAAAGCTATGTAACCTTCAATTTCCATAACCAA 5640
Db 870 AGGAGGTTTACCTCTCATTCTCTAGAGAAAAGCTATGTAACCTTCAATTTCCATAACCAA 929
Qy 5641 TACTTTATATATGTAAGTTTATTTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700
Db 930 TATTTTATATGTAAGTTTATTTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 989
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Db 1049 ATTTATGATTAATTAATTAAGAAAACAGATATCTTAGGCTTTTAAATAACACATGAATATCA 1108
Qy 5821 TAAA 5824
Db 1109 TAAA 1112

RESULT 10
US-10-627-273-7
; Sequence 7, Application US/10627273
; Publication No. US20040110189A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; APPLICANT: Renauld, Jean-Christophe
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; TITLE OF INVENTION: (Tifs) The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5543.2
; CURRENT APPLICATION NUMBER: US/10/627,273
; CURRENT FILING DATE: 2003-07-25
; PRIOR FILING DATE: US/09/751,797
; PRIOR FILING DATE: 2000-12-29
; PRIOR APPLICATION NUMBER: 09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 29
; SEQ ID NO 7
; LENGTH: 1119
; TYPE: DNA
; ORGANISM: Mus musculus
; FEATURE:
US-10-627-273-7

Query Match 9.4%; Score 555.2; DB 7; Length 1119;
Best Local Similarity 96.0%; Pred. No. 8.9e-113; Indels 1; Gaps . 1;
Matches 580; Conservative 0; Mismatches 23;

Qy 5221 ATAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCTGTTTATGTC 5280
Db 510 AAAGCTTGGAGAGATCGGAGAGATCAAGCGGATCGGGAACTGGACCTGCTGTTTATGTC 569
Qy 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAGAACTGCTCTCTTCCT 5340
Db 570 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAGAACTGCTCTCTTCCT 629
Qy 5341 GCCTTCTAAAAGAACATAGATCCCTGAATGGACTTTTCTTAAAGGAAAGTGAGAA 5400
Db 630 GCCTTCTAAAAGAACATAGATCCCTGAATGGACTTTTCTTAAAGGAAAGTGAGAA 689
Qy 5401 GCTAACGCTCCACCATCATTTAGAAATTTTACATGAAACCTGGCTCAGTTGAAAGAGAAA 5460
Db 593 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAGAACTGCTCTCTTCCT 652
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Db 690 GCTAACGCTCCATCATCATTAGAAAGATTTTCAATGAAACCTGGCTCAGTTGAAAAGAAAA 749
Qy 5461 TAGTGTCAGGTTGTCATGAGACGAGGTTAGACTTTGATAAACCAAGAGATTCATTGACA 5520
Db 750 TAGTGTCAGGTTGTCATGAGACGAGGTTAGACTTTGATAAACCAAGAGATTCATTGACA 809
Qy 5521 ATATTTTATTTGTCATGATTAATGCAACAGAAAAGTATGACTTTTAAAAAATTGTTTGA 5580
Db 810 ATATTTTATTTGTCATGATTAATGCAACAGAAAAGTATGACTTTTAAAAAATTGTTTGA 869
Qy 5581 AGGAGGTTTACCTCTCATTCTCTAGAGAAAAGCTATGTAACCTTCAATTTCCATAACCAA 5640
Db 870 AGGAGGTTTACCTCTCATTCTCTAGAGAAAAGCTATGTAACCTTCAATTTCCATAACCAA 929
Qy 5641 TACTTTATATATGTAAGTTTATTTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700
Db 930 TATTTTATATGTAAGTTTATTTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 989
Qy 5701 ATATGGAATTTATTTATAGAAAATTTATCTGATGTTGATATTTAGATATAAGCAATAAT 5760
Db 990 ATATGGAATTTATTTATAGAAAATTTATCTGATGTTGATATTTAGATATAAGCAATAAT 1048
Qy 5761 ATTTATGATTAATTAATTAAGAAAACAGATATCTTAGGCTTTTAAATAACACATGAATATCA 5820
Db 1049 ATTTATGATTAATTAATTAAGAAAACAGATATCTTAGGCTTTTAAATAACACATGAATATCA 1108
Qy 5821 TAAA 5824
Db 1109 TAAA 1112

RESULT 11
US-10-084-298-3
; Sequence 3, Application US/10084298
; Publication No. US20030099649A1
; GENERAL INFORMATION:
; APPLICANT: Jacobs, Kenneth
; APPLICANT: Pittman, Debra
; APPLICANT: Fouser, Lynette
; APPLICANT: Spaulding, Vikki
; APPLICANT: Xuan, Dejun
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory
; TITLE OF INVENTION: Disorders
; FILE REFERENCE: GI5358 CIP
; CURRENT APPLICATION NUMBER: US/10/084,298
; CURRENT FILING DATE: 2002-09-10
; PRIOR FILING DATE: 60/270,823
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/281,353
; PRIOR FILING DATE: 2001-04-03
; PRIOR APPLICATION NUMBER: 60/131,473
; PRIOR FILING DATE: 1999-04-28
; PRIOR APPLICATION NUMBER: 09/561,811
; PRIOR FILING DATE: 2000-04-28
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 1166
; TYPE: DNA
; ORGANISM: Murine
US-10-084-298-3

Query Match 9.4%; Score 555.2; DB 5; Length 1166;
Best Local Similarity 96.0%; Pred. No. 9.1e-113; Indels 1; Gaps 1;
Matches 580; Conservative 0; Mismatches 23;

Qy 5221 ATAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCTGTTTATGTC 5280
Db 533 AAAGCTTGGAGAGATCGGAGAGATCAAGCGGATCGGGAACTGGACCTGCTGTTTATGTC 592
Qy 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAGAACTGCTCTCTTCCT 5340
Db 593 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAGCTAGAAAACGAGAACTGCTCTCTTCCT 652
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[illegible]

RESULT 14  
US-11-157-387-3  
; Sequence 3, Application US/11157387  
; Publication No. US20050238648A1  
; GENERAL INFORMATION:  
; APPLICANT: Jacobs, Kenneth  
; APPLICANT: Pittman, Debra  
; APPLICANT: Fouser, Lynette  
; APPLICANT: Spaulding, Vikki  
; APPLICANT: xuan, Dejun  
; TITLE OF INVENTION: Composition and Method for Treating Inflammatory  
; TITLE OF INVENTION: Disorders  
; FILE REFERENCE: G15358 CIP  
; CURRENT APPLICATION NUMBER: US/11/157,387  
; CURRENT FILING DATE: 2005-06-20  
; PRIOR APPLICATION NUMBER: US/10/084,298  
; PRIOR FILING DATE: 2002-09-10  
; PRIOR APPLICATION NUMBER: 60/270,823  
; PRIOR FILING DATE: 2001-02-23

;	PRIOR APPLICATION NUMBER: 60/281,353	
;	PRIOR FILING DATE: 2001-04-03	
;	PRIOR APPLICATION NUMBER: 60/131,473	
;	PRIOR FILING DATE: 1999-04-28	
;	PRIOR APPLICATION NUMBER: 09/561,811	
;	PRIOR FILING DATE: 2000-04-28	
;	NUMBER OF SEQ ID NOS: 10	
;	SOFTWARE: PatentIn Ver. 2.1	
;	SEQ ID NO 3	
;	LENGTH: 1166	
;	TYPE: DNA	
;	ORGANISM: Murine	
US-11-157-387-3		
	Query Match 9.4%; Score 555.2; DB 10; Length 1166;	
	Best Local Similarity 96.0%; Pred.No. 9.1e-113;	
	Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;	
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Qy	5281 TCTGAGAAATGCTTGGCTCTGACGAGAGAAGAACTAGAAAACGAAGAACTGCTCCTTCCT	5340
Db	593 TCTGAGAAATGCTTGGCTCTGACGAGAGAAGAACTAGAAAACGAAGAACTGCTCCTTCCT	652
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Qy	5461 TAGTGCAAGTTGTCATGACAGACAGAGGTAGACTTGTATACCACCAAGAAATTCATTGACA	5520
Db	773 TAGTGCAAGTTGTCATGACAGACAGAGGTAGACTTGTATACCACCAAGAAATTCATTGACA	832
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Qy	5581 AGGAGGTTACCTCTCATTTCTTAGAAGAAAAAGCCTATGTAATTCATTTCCATAACCAA	5640
Db	893 AGGAGGTTACCTCTCATTTCTTAGAAGAAAAAGCCTATGTAATTCATTTCCATAACCAA	952
Qy	5641 TACTTTATATATGTAAGTTTATTTATATAGTATACATTTTATTTATGTCAGTTTATTA	5700
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Db	1013 ATATGGATTTATTTATAGAAAAATTAATCTGCTATTTGATATTTAGTATAGGCCAATAAT	1071
Qy	5761 ATTTATGATAATAACTATAGAAAAACAGATATCTTAGGCTTTTAAATAACACATGAATATCA	5820
Db	1072 ATTTATGACAAATAACTATAGAAAAACAGATATCTTAGGCTTTTAAATAACACATGAATATCA	1131
Qy	5821 TAAA 5824	
Db	1132 TAAA 1135	

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RESULT 15
US-10-090-365-40
; Sequence 40, Application US/10090365
; Publication No. US2003007706A1
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
; APPLICANT: Xu, Wenfeng
; APPLICANT: Kindsvogel, Wayne
; APPLICANT: Chen, Zhi
; TITLE OF INVENTION: Mouse Cytokine Receptor

```



; FILE REFERENCE: 01-08  
; CURRENT APPLICATION NUMBER: US/10/090,365  
; CURRENT FILING DATE: 2002-03-04  
; PRIOR APPLICATION NUMBER: US 60/273,035  
; PRIOR FILING DATE: 2001-03-02  
; PRIOR APPLICATION NUMBER: US 60/279,232  
; PRIOR FILING DATE: 2001-03-27  
; NUMBER OF SEQ ID NOS: 49  
; SOFTWARE: FastSeq for Windows Version 3.0  
; SEQ ID NO 40  
; LENGTH: 1050  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (50)...(589)  
US-10-090-365-40

Query Match 9.1%; Score 541.4; DB 5; Length 1050;  
Best Local Similarity 99.8%; Pred. No. 1e-109;  
Matches 542; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
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Db 508 AAAGCTTGGAGAGCGGAGAGATCAAGCGATCGGGGAACTGGACCTGCTGTTTATGTC 567  
Qy 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAGCTAGAAAACGAAAGAACTGCTCCTTCCT 5340  
Db 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAGCTAGAAAACGAAAGAACTGCTCCTTCCT 627  
Qy 5341 GCCTTCTAAAAGAAACAATAGATCCCTGAATGGACTTTTTTACTAAAGGAAAAGTGAGAA 5400  
Db 628 GCCTTCTAAAAGAAACAATAGATCCCTGAATGGACTTTTTTACTAAAGGAAAAGTGAGAA 687  
Qy 5401 GCTAAAGCTCCACCATCATTTAGAGATTTTCACATGAACCTGGCTCAGTTCAAAGAGNAAA 5460  
Db 688 GCTAAAGCTCCACCATCATTTAGAGATTTTCACATGAACCTGGCTCAGTTCAAAGAGNAAA 747  
Qy 5461 TAGTGTCAGGTGTGTCATGAGACAGAGGTAGACTTGATTAACCAAGAGATTCAATTGACA 5520  
Db 748 TAGTGTCAGGTGTGTCATGAGACAGAGGTAGACTTGATTAACCAAGAGATTCAATTGACA 807  
Qy 5521 ATATTTTATTGTCATTGATTAATGCAACAGAAAAGTATGTACTTTAAAAAATTGTTTGA 5580  
Db 808 ATATTTTATTGTCATTGATTAATGCAACAGAAAAGTATGTACTTTAAAAAATTGTTTGA 867  
Qy 5581 AGGAGGTACCTCTCATCTCTAGAGAAAAGCCTATGTAATCTTCAATTTCCATAACCAA 5640  
Db 868 AGGAGGTACCTCTCATCTCTAGAGAAAAGCCTATGTAATCTTCAATTTCCATAACCAA 927  
Qy 5641 TACTTTATATATGTAAGTATTATTATTATTAAGTATACATTTTATTATGTCAGTTTATTA 5700  
Db 928 TACTTTATATATGTAAGTATTATTATTATTAAGTATACATTTTATTATGTCAGTTTATTA 987  
Qy 5701 ATATGGATTTATTATAGAAAAATTAFTCTGATGTTGATATTTGAGTATAAGCAAAATAAT 5760  
Db 988 ATATGGATTTATTATAGAAAAATTAFTCTGATGTTGATATTTGAGTATAAGCAAAATAAT 1047  
Qy 5761 ATT 5763  
Db 1048 ATT 1050

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GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: December 20, 2005, 13:17:12 ; Search time 413.821 Seconds  
(without alignments)  
7442.822 Million cell updates/sec

Title: US-09-751-797-29

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Gapop 10.0 , Gapext 1.0

Searched: 4168288 seqs, 259477437 residues

Total number of hits satisfying chosen parameters: 8336576

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

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8: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq2.\*  
9: /cgn2\_6/ptodata/2/pubpna/US11\_NEW\_PUB.seq3.\*  
10: /cgn2\_6/ptodata/2/pubpna/US60\_NEW\_PUB.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	5935	100.0	5935	7 US-11-177-987-42	Sequence 42, Appl
2	4245.2	71.5	7445	7 US-11-177-987-8	Sequence 8, Appl
3	650	11.0	4797	7 US-11-177-987-26	Sequence 26, Appl
4	602.4	10.1	1111	7 US-11-177-987-9	Sequence 9, Appl
5	555.2	9.4	1119	7 US-11-177-987-7	Sequence 7, Appl
6	215	3.6	1152	7 US-11-102-240-153	Sequence 153, App
7	159.4	2.7	187745	7 US-11-121-086-83	Sequence 83, Appl
8	145	2.4	171936	7 US-10-933-025-24	Sequence 24, Appl
9	142	2.4	260209	6 US-10-933-025-23	Sequence 23, Appl
10	139	2.3	17004	7 US-11-176-253-1	Sequence 1, Appl
11	133	2.2	135019	6 US-10-849-438-11	Sequence 11, Appl
12	128.8	2.2	10166	7 US-11-147-606-3	Sequence 3, Appl
13	127.6	2.1	690	7 US-11-177-987-25	Sequence 25, Appl
14	126.4	2.1	126552	7 US-11-121-086-1	Sequence 1, Appl
15	125.4	2.1	2695	7 US-11-000-365-35	Sequence 35, Appl
16	125.4	2.1	2695	7 US-11-032-794-35	Sequence 24, Appl
17	124.6	2.1	171936	6 US-10-933-025-24	Sequence 2, Appl
18	120.4	2.0	4621	6 US-10-960-644-2	Sequence 3, Appl
19	119.6	2.0	10166	7 US-11-147-606-3	Sequence 18, Appl
20	117.2	2.0	418	7 US-11-177-987-18	Sequence 18, Appl
21	117	2.0	184868	7 US-11-121-086-88	Sequence 88, Appl
22	114.8	1.9	2868	9 US-11-012-762-5	Sequence 5, Appl
23	111.4	1.9	394468	6 US-10-995-561-13473	Sequence 13473, A

RESULT 1

US-11-177-987-42

; Sequence 42, Application US/11177987

; Publication No. US20050271619A1

; GENERAL INFORMATION:

; APPLICANT: Dumoutier, Laure

; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac

; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof

; FILE REFERENCE: LUD 5664

; CURRENT FILING DATE: 2005-07-07

; PRIOR APPLICATION NUMBER: US/11/177,987

; PRIOR FILING DATE: 2000-07-27

; PRIOR APPLICATION NUMBER: US09/419,568

; PRIOR FILING DATE: 1999-10-18

; PRIOR APPLICATION NUMBER: US09/354,243

; PRIOR FILING DATE: 1999-07-16

; PRIOR APPLICATION NUMBER: US09/178,973

; PRIOR FILING DATE: 1998-10-26

; NUMBER OF SEQ ID NOS: 43

; SEQ ID NO 42

; LENGTH: 5935

; TYPE: DNA

; ORGANISM: Mus musculus

; FEATURE:

US-11-177-987-42

Query Match 100.0%; Score 5935; DB 7; Length 5935;

Best Local Similarity 100.0%; Pred. No. 0;

Matches 5935; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1 GAATTCAGTCCACATGCAATCAATCGAATACCTTTGTAATTTCTTCTTCAAAATATCC 60

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Db 61 ATCTATATAGTATAAGTTATTTAGGATCAATTTAAATAATGTTTGGAGACTTATGTTT 120

Qy 121 GCACAGTAAATGTCAGAGAGAAATAGCAAAATGATAGTATTATTTTAAAAAAT 180

Db 121 GCACAGTAAATGTCAGAGAGAAATAGCAAAATGATAGTATTATTTTAAAAAAT 180

Qy 181 CTATGCTAAAAATGCTTATTAGATTGTTCACTGCAATTTCCAAACTTAACCTTGACCT 240

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Db 181 CTATGCTAAAAATGCTCTATTAGATTGTTCACTACTAGACATTTCCAAAATTTAACTTGACCT 240  
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Db 241 TGGCTATGATTTCAACCTTTGTTATTTGCACTTACCATAAATGTTGCTCTCACTTACCATGC 300  
Qy 301 TATCCGACGAGCATGTTCCCTCGATGTTTTCCTTTGCTCTCTCGCTAAACAGGCTCTC 360  
Db 301 TATCCGACGAGCATGTTCCCTCGATGTTTTCCTTTGCTCTCTCGCTAAACAGGCTCTC 360  
Qy 361 CTCTCAGTTATCAAACTTTTGACACTTGTGGCATCGGTGATGGCTGTCCTCGAGAAAATCTA 420  
Db 361 CTCTCAGTTATCAAACTTTTGACACTTGTGGCATCGGTGATGGCTGTCCTCGAGAAAATCTA 420  
Qy 421 TGAGTTTTTCCCTTATGGGACCTTTGGCCGACCTGCTCTCATTTGGCCCTGTGGG 480  
Db 421 TGAGTTTTTCCCTTATGGGACCTTTGGCCGACCTGCTCTCATTTGGCCCTGTGGG 480  
Qy 481 CCCAGGAGGCAAAATGCGCTGCCCATCAACCCCGGTGCAAGCTTGAGGTGCCAACTTCC 540  
Db 481 CCCAGGAGGCAAAATGCGCTGCCCATCAACCCCGGTGCAAGCTTGAGGTGCCAACTTCC 540  
Qy 541 AGCAGCCGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGGAGGTACAGCTGCATCTCT 600  
Db 541 AGCAGCCGTACATCGTCAACCGCACCTTTATGCTGGCCAAAGGAGGTACAGCTGCATCTCT 600  
Qy 601 TTCTCTCATACCGCTTGCCATTTCTCTGAAGCACTTGCAAAATCTTTTAAAGGCGGCTTT 660  
Db 601 TTCTCTCATACCGCTTGCCATTTCTCTGAAGCACTTGCAAAATCTTTTAAAGGCGGCTTT 660  
Qy 661 ATCTCCGAGGTCTCACTACCTATGTTTCTGCTCTTTTAGAGACTCTTTTAAAGACTGGA 720  
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Qy 721 TCTTTTCTATTTCTATTTCAAGGTCTCAGGACCAATTTCTATCTGGCCCTTCAGGACAC 780  
Db 721 TCTTTTCTATTTCTATTTCAAGGTCTCAGGACCAATTTCTATCTGGCCCTTCAGGACAC 780  
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Db 901 GGACATACATCTCACTTACAGGCTTTTCTTCATCTCTCTTGTCAACCCAGGCACTTAGGG 960  
Qy 961 TTTTCTCTTTTCAGGCCAGCTTGCAGATAAACAACAACAGAGCTCGGCTCATCGGGAG 1020  
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Qy 1141 GCTAAGGATCAGTGTACTGATGAGCAGGTGCTCAACTTTCACCCCTGGAGACATTTCTG 1200  
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Qy 1201 CTCCCCAGTCAGACAGGTTCCGGCCCTACATGACGAGGTTGGTCCCTTTCTGACCAAA 1260  
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Db 1381 AGGCGCTCAGCACCAACCATCATAGGCCACTGTGAATAAGGTCAAAAAGGCTTTGGCTT 1440  
Qy 1441 CAATTGAGTAAATCTTTGAGTTTGTATTAGTTAAGCTTTTATTGTTTTTATCCATGGAAAG 1500  
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Db 1681 TGGCACAGACAAAGTTGAAAGGAAAGAGAGATGGAGAGGCTCAATGTTGGGGGTG 1740  
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Qy 1861 GGCTTTTTCACACGAGAAATTTTATGCTCATCTCTTGTGCTACACTCCACACTTTTGTAGTGA 1920  
Db 1861 GGCTTTTTCACACGAGAAATTTTATGCTCATCTCTTGTGCTACACTCCACACTTTTGTAGTGA 1920  
Qy 1921 GTTAAGCTCAGGTTTCGTTTCTACCGTTCTTGTCTACCTGGTGAACCTTTCAGTAGGATTC 1980  
Db 1921 GTTAAGCTCAGGTTTCGTTTCTACCGTTCTTGTCTACCTGGTGAACCTTTCAGTAGGATTC 1980  
Qy 1981 CCAGAAGCAGGACAGCTCTTCTGTAGGAGGAGCCTGGATTTTCAGTGTCTCTAGAGAAC 2040  
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Qy 2041 GAAATAGCTCAGAGAAATCTAGGTCAACGTGAAATCTAGGTCAACGCGGGCAAAAATGACT 2100  
Db 2041 GAAATAGCTCAGAGAAATCTAGGTCAACGTGAAATCTAGGTCAACGCGGGCAAAAATGACT 2100  
Qy 2101 GAAAGCTCTATTTCCAGGTGAAACGCTCAACGCTGCTCAGATATCTAGGATTTGGGCTTC 2160  
Db 2101 GAAAGCTCTATTTCCAGGTGAAACGCTCAACGCTGCTCAGATATCTAGGATTTGGGCTTC 2160  
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Qy 2281 AAGCCACAATACTAAGCCATTCAGTAGGAGACGTGGGATTTCTTTCTCTCTCCCACT 2340  
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Qy	2401	CTGCACCTGCATCTAGCTGGGTCTATAGATCTTTCAATCTGTGTCATAAATTTGTAAAGTCA	2460
Db	2401	CTGCACCTGCATCTAGCTGGGTCTATAGATCTTTCAATCTGTGTCATAAATTTGTAAAGTCA	2460
Qy	2461	CAATTCCTGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTCATGAGCACTTGCCTCGGAGGA	2520
Db	2461	CAATTCCTGGAGCTAGCAGAAAGCTTAGCTCAGCCAGTCTCATGAGCACTTGCCTCGGAGGA	2520
Qy	2521	TGGCTTGTGACAGAGTCAATGTCTAGAAAGACAGCATCCCTGATTTCCCAAGCTCTGCACCTTGC	2580
Db	2521	TGGCTTGTGACAGAGTCAATGTCTAGAAAGACAGCATCCCTGATTTCCCAAGCTCTGCACCTTGC	2580
Qy	2581	CTAGTGGCCAGTGTAAATTACTTTAGCCTGATTAAGTATTGGGAAAGCCAAATTTCCCAACC	2640
Db	2581	CTAGTGGCCAGTGTAAATTACTTTAGCCTGATTAAGTATTGGGAAAGCCAAATTTCCCAACC	2640
Qy	2641	GACCTACATAATCCGAAGAAGCATGCAATTCGAAACTAGAAAGCTGGGCAACAACCTTACTA	2700
Db	2641	GACCTACATAATCCGAAGAAGCATGCAATTCGAAACTAGAAAGCTGGGCAACAACCTTACTA	2700
Qy	2701	GAGATGATTTTTGAGCTCATTTAAACTCATGCTCTGAAATGTGATCAAAATCAAACCCAGAAT	2760
Db	2701	GAGATGATTTTTGAGCTCATTTAAACTCATGCTCTGAAATGTGATCAAAATCAAACCCAGAAT	2760
Qy	2761	AACAACAAAAGAGCTGGATTTGCAAAATAGACAAGTATTTAGAAATCACTGGTATTAACAG	2820
Db	2761	AACAACAAAAGAGCTGGATTTGCAAAATAGACAAGTATTTAGAAATCACTGGTATTAACAG	2820
Qy	2821	CTGTCACTTTAATTAATAATAGTGTCTATTTAGCTGCCTATTTAGATTTAAACACAAGA	2880
Db	2821	CTGTCACTTTAATTAATAATAGTGTCTATTTAGCTGCCTATTTAGATTTAAACACAAGA	2880
Qy	2881	GTGGATAAATCTTCCCAATTTTACTGGGCTGGTTTCAATAGAGTAAAAATATCAGTCAATAGA	2940
Db	2881	GTGGATAAATCTTCCCAATTTTACTGGGCTGGTTTCAATAGAGTAAAAATATCAGTCAATAGA	2940
Qy	2941	TTAATTAATAGTGATCAAGAAAGTATGAGTTGGAAACCCCTTTCTTTTACCTTTTACCTTCAT	3000
Db	2941	TTAATTAATAGTGATCAAGAAAGTATGAGTTGGAAACCCCTTTCTTTTACCTTTTACCTTCAT	3000
Qy	3001	TCCTTAGTTATATTTTTTTTTTTCTTCACACCTGATCAAGCCACTAGTAAAGCACTATCTG	3060
Db	3001	TCCTTAGTTATATTTTTTTTTTTCTTCACACCTGATCAAGCCACTAGTAAAGCACTATCTG	3060
Qy	3061	CTCGAGCTATATATGACTTTACAGCAAAACAACATTCGTGTGGGCTCTTTGGGGAAG	3120
Db	3061	CTCGAGCTATATATGACTTTACAGCAAAACAACATTCGTGTGGGCTCTTTGGGGAAG	3120
Qy	3121	GGAAACGAGTAGCAGGAGGCTCAGGCTAGCAAGTCTGGAATCAACCTTAAAGCCAGAGGCA	3180
Db	3121	GGAAACGAGTAGCAGGAGGCTCAGGCTAGCAAGTCTGGAATCAACCTTAAAGCCAGAGGCA	3180
Qy	3181	TGGTTGATAGCAGAAAGTGAAGCTCTTACAAAGTGGGTGCTTAAGTAAATCAGAAAC	3240
Db	3181	TGGTTGATAGCAGAAAGTGAAGCTCTTACAAAGTGGGTGCTTAAGTAAATCAGAAAC	3240
Qy	3241	AGGAAGGCTCTGGTTGATGGAAATTATCAGTAAGATATCTACCCCTTATCTCCTTCTTCTAT	3300
Db	3241	AGGAAGGCTCTGGTTGATGGAAATTATCAGTAAGATATCTACCCCTTATCTCCTTCTTCTAT	3300
Qy	3301	AGAAGCTTAAACCGTCTCTCCTTCTGTGTGTAGGCTGATAAACAACGCTGTTTTCTTTTG	3360
Db	3301	AGAAGCTTAAACCGTCTCTCCTTCTGTGTGTAGGCTGATAAACAACGCTGTTTTCTTTTG	3360
Qy	3361	AGTGTTCAATGGCTTTGACAGATTTTTCAGTGTCTCTGCCAGTTCTTGTAGAGGTTTGTTC	3420
Db	3361	AGTGTTCAATGGCTTTGACAGATTTTTCAGTGTCTCTGCCAGTTCTTGTAGAGGTTTGTTC	3420
Qy	3421	CTTTCACACCTGGGCTCGATGCTTAGCATGCCAAGGACACACCTTCTGAAATGCTGTGTA	3480
Db	3421	CTTTCACACCTGGGCTCGATGCTTAGCATGCCAAGGACACACCTTCTGAAATGCTGTGTA	3480
Qy	3481	AAAGGTTATTATTCAATTTACTTTGTCTTTTGGAAAGGTGAAGTGTGTGTGAGAAAGAACCTC	3540

[illegible]



QY 652 GGGCGCTTTATCTCCGAGGTCTCACTACCTATGTTTTTCTGTCTCTTTAGAGACTCTTTA 711  
DB 2331 GGGCGCTTTATCTCCGAGGTCTCACTACCTATGTTTTTCTGTCTCTTTAGAGACTCTTTA 2390  
QY 712 AGAGTGGATCTTTTCTATTTCTATTTCAAGGTCTCAGACCAATTTCCCTATCTTGGCCT 771  
DB 2391 AGAGTGGGTCTTTTCTATTTCTATTTCAAGGTCTCAGACCAATTTCCCTATCTTGGCCT 2450  
QY 772 TCAGGACACATATACTCAATTTTTATCTACAGAGCGGTTTTAGAAAGCCACCCACGACTG 831  
DB 2451 TCAGGACACATATACTCAATTTTTATCTACAGAGCGGTTTTAGAAAGCCACCCACGACTG 2510  
QY 832 CAATACCTTCCATCTCTGTGTCTCTTTCTGAACCTCATACTCTCTTGGTCTACTCTCTGAG 891  
DB 2511 CAATACCTTCCATCTCTGTGTCTCTTTCTGAACCTCATACTCTCTTGGTCTACTCTCTGAG 2570  
QY 892 ACCCACTGGGACATACATCTCTACTTTACAGGCTTTTCTCCATCTCTTGTCACCGAG 951  
DB 2571 ACCCACTGGGACATACATCTCTACTTTACAGGCTTTTCTCCATCTCTTGTCACCGAG 2630  
QY 952 CACTTAGGGTTTTCTCTCTTTTCAGGCCAGCTTGCAGATAACAACACAGACGTCGGGCTC 1011  
DB 2631 CACTTAGGGTTTTCTCTCTTTTCAGGCCAGCTTGCAGATHAACACAGACGTCGGGCTC 2690  
QY 1012 ATCGGGAGAACTGTTCCGAGGAGTCAGTGTAACTCTCTCACTGTGTATGACGAGGCTAG 1071  
DB 2691 ATCGGGAGAACTGTTCCGAGGAGTCAGTGTAACTCTCTCACTGTGTATGACGAGGCTAG 2750  
QY 1072 CTGGGAGCTGTGGACCTCTGGGATAGTCTGAAGTATGACCCCTGTCTCTTGTGTC 1131  
DB 2751 CTGGGAGCTGTGGACCTCTGGGATAGTCTGAAGTATGACCCCTGTCTCTTGTGTC 2810  
QY 1132 TACTCGAGCTAAGGATCAGTCTACTCTGATGAGCAGGTCTCACTTCACTTCACTTGGAA 1191  
DB 2811 TACTCGAGCTAAGGATCAGTCTACTCTGATGAGCAGGTCTCACTTCACTTCACTTGGAA 2870  
QY 1192 GACATCTCTCTCCCTCAGTCTCAGACAGGTTCCGGCCCTTACATGACGAGGAGTGGTCTTC 1251  
DB 2871 GACATCTCTCTCCCTCAGTCTCAGACAGGTTCCGGCCCTTACATGACGAGGAGTGGTCTTC 2930  
QY 1252 CTGACCAAACTCAGCAATCAGCTCAGTCTCTGTGTAAAGTCTGGTCTGGTCTACTATGCT 1311  
DB 2931 CTGACCAAACTCAGCAATCAGCTCAGTCTCTGTGTAAAGTCTGGTCTGGTCTACTATGCT 2990  
QY 1312 CCTCTCTCTCTCTCTTCTATTTCCAGTAAGAACCCGAGGCTCTGCCCTCTCTCTCTTCA 1371  
DB 2991 CCTCTCTCTCTCTCTTCTATTTCCAGTAAGAACCCGAGGCTCTGCCCTCTCTCTCTTCA 3050  
QY 1372 AGAGTGAGGAGGGCTCAGCACCAACCATCATAGGCCACTTGAATAGGTCAACAAG 1431  
DB 3051 AGAGTGAGGAGGGCTCAGCACCAACCATCATAGGCCACTTGAATAGGTCAACAAG 3110  
QY 1432 CTTTGGCTTCAATTTAGTAATATCTTTGAGTTGTTATTTAGTTTATTTTATTTATC 1491  
DB 3111 CTTTGGCTTCAATTTAGTAATATCTTTGAGTTGTTATTTAGTTTATTTTATTTATC 3170  
QY 1492 CATGGAAGAAATCAACTCAATTTCTGATGATGAGAAAGTGTGGGAACGAAAGAAG 1551  
DB 3171 CATGGAAGAAATCAACTCAATTTCTGATGATGAGAAAGTGTGGGAACGAAAGAAG 3230  
QY 1552 CCTAGATAGAGAAACAGATCTGCTGATACAGTACTTTATGGGGGGGGGGGGGGGGGG 1611  
DB 3231 CCTAGATAGAGAAACAGATCTGCTGATATAGTACTTAT ---GGGGGGGAGCGGGGGCG 3286  
QY 1612 ATATCCACTGAGTCAAGTACTTTTGGGAGAGAAATCCACTGAGTCAAGTACTTGT -- 1669  
DB 3287 ATATCCACTGAGTCAAGTACTTTTGGGAGAGAAATCCACTGAGTCAAGTACTTGTG 3346  
QY 1670 -----GGGGGAAGGAATGGCAGCAGGCAAAAGT 1697  
DB 3347 GCATGGAGATCCACTGAGTCAAGTACTTTGGGGGGGGGGGAATGGCAGCAGGCAAAAGT 3406

QY 1698 TGAAGGGAAGAGAGATGGAGAGCCCTCAATGTTGGGGGTGTGAAAGGTCACTCCTTT 1757  
DB 3407 TGAAGGGA ---AGGAAGATGGAGAGCCCTCATGCTTGGGGGTGTGAAAGGTCACTCC -TT 3462  
QY 1758 TTCCATGTGATGAGAGATTAAGAAATCAGTGTGTGAGTTTGTATGTTCTTTCAGACACCCC 1817  
DB 3463 TTCCATGTGATGAGAGATTAAGAAACCAAGTGTGTGAGTTTGTATGTTCTTTCAGACACCCC 3522  
QY 1818 AA-----CTATGGCAGACTGTGGGAGACCTTGGCAATTTAGGGAA 1855  
DB 3523 CAACTATGAACATATCCAGGAGGGGCGAGACTGTGGGAGACCTTGGCAATTTAGGGAA 3582  
QY 1856 GGGCGGCTTTTTCACAGAGAACTTTATGCTCATCTCTTGTGCTACACTCCACCTTTG 1915  
DB 3583 GGGCGGCTTTTTCACAGAGAACTTTATGCTCATCTCTTGTGCTACACTCCACCTTTG 3642  
QY 1916 ATCAGGTTAAGCTCAGGTTTCTACCGTTTCTGCTACTGCTGTAATCTTTCAGTAGG 1975  
DB 3643 ATCAGGTTAAGCTCAGGTTTCTACCGTTTCTGCTACTGCTGTAATCTTTCAGTAGG 3702  
QY 1976 ATTCCCAAGAGCAGGACAGCTCTTCTGTAAAGGAGGACCTTGGATTTTCAGTGTCTAG 2035  
DB 3703 ATTCCCAAGAGCAGGACAGCTCTTCTGTAAAGGAGGACCTTGGATTTTCAGTGTCTAG 3762  
QY 2036 AGAACGAATATGCTCAGAGATCTAGGTCACCTGTAATCTAGGTCACAGGGGCAAAA 2095  
DB 3763 AGAACGAATATGCTCAGAGATCTAGGTCACCTGTAATCTAGGTCACAGGGGCAAAA 3822  
QY 2096 TGAAGTAAAGCTCTTATTCAGGTCAGGTCAGGTCAGGTCAGATATCTTTCAGTGTATGG 2155  
DB 3823 TGAAGTAAAGCTCTTATTCAGGTCAGGTCAGGTCAGGTCAGATATCTTTCAGTGTATGG 3882  
QY 2156 GCTCCCAAGGATTAAGATCTGTTAGTCTGCTCTTCTTATTTTCAGGACATCAGTGGT 2215  
DB 3883 GCTCCCAAGGATTAAGATCTGTTAGTCTGCTCTTCTTATTTTCAGGACATCAGCGGT 3942  
QY 2216 GACGACAGAAATCCAGAGAAATGTCAGAGGCTGAAGGACAGTGAAGGATCTA 2275  
DB 3943 GACGACAGAAATCCAGAGAAATGTCAGAGGCTGAAGGACAGTGAAGGATCTA 4002  
QY 2276 TTGGCAAGCCACAATTAAGCCATTCAGTAGGAGCGTGGGATTTCTTCTCTGCTTC 2335  
DB 4003 TTGGCAAGCCACAATTAAGCCATTCAGTAGGAGCGTGGGATTTCTTCTCTGCTTC 4062  
QY 2336 CCACTCTCTCTACTTTGTAAACATTTCTTGTACTGTCTACTGTCTGGTCCATTACTCA 2395  
DB 4063 CCACTCTCTCTACTTTGTAAACATTTCTTGTACTGTCTACTGTCTGGTCCATTACTCG 4122  
QY 2396 CTTAGCTGCACCTGCATCTAGCTGGGTCTATAGATCTTTCAATCTGTGTCTAAATTTGTA 2455  
DB 4123 CTTAGCTGCACCTGTATCTAGCTGGGTCTATAGATCTTTCAATCTGTGTCTAAATTTGTA 4182  
QY 2456 AGTCACAAATCTGGAGTAGCAGAAAGCTTAGCTCAGCCAGTCTCATGACACTTGTCTCG 2515  
DB 4183 AGTCACAAATCTGGAGTAGCAGAAAGCTTAGCTCAGCCAGTCTCATGACACTTGTCTCG 4242  
QY 2516 GAGGATGGCTTGTGACAGAGTCAATGCTAGAAGACAGATCCCTGATTTCCAGCTCTGCA 2575  
DB 4243 GAGGATGGCTTGTGACAGAGTCAATGCTAGAAGACAGATCCCTGATTTCCAGCTCTGCA 4302  
QY 2576 CTTGCTCTAGTGGCCACGCTGTAATTAATTTAGCTGTATTAAGTATTTGGGAAGCAATTC 2635  
DB 4303 CTTGCTCTAGTGGCCACGCTGTAATTAATTTAGCTGTATTAAGTATTTGGGAAGCAATTC 4362  
QY 2636 CCAAGGACCTACATATCCGAAGACGATGCTATGAAGCTAGAAAGCTGGGCAAACT 2695  
DB 4363 CCAAGGACCTACATATCTGAAGACGATGCTATGAAGCTAGAAAGCTGGGCAAACT 4422  
QY 2696 TACTAGAGATGATTTTGTAGCTCAATTAAGTGTCTTCAAAATGTGATCAAAATCAACCC 2755  
DB 4423 TACTAGAGATGATTTTGTAGCTCAATTAAGTGTCTTCAAAATGTGATCAAAATCAACCC 4482  
QY 2756 AGAATAACCAAAAGAGCTGGATTTGCAATATAGGCAAGTATTTTAGAATCACTGGTATT 2815



Db		4483	AGATAACAACAAGAGCTGGATTTGCAATAGGACAAGTATTTAGAACTACCTGGTATT	4542
Qy		2816	AACAGCTGTCATCTTAATAATAATATAGTGTCTATTTAGCTGCGCTATTTAAGATTAACA	2875
Db		4543	AATAGCTATCATCTTAATAATAATATAGGGCTATATA----TATATTTAAGATTAACA	4598
Qy		2876	CAAGAGTGGATACTTCCCAATTTACTTGGGCGTGGTTTCAATAGAGTAAATAATACAGTC	2935
Db		4599	CAAGAGTGGATAGCCTCCCAATTTACTTGGGCGTGGTTTCAAAAGAGTAAATAATACAGTC	4658
Qy		2936	ATAGATTAATATATAGTGTCAATGAAGTATGAGTTGGAAACCCCTTTCCTTACTTTTACCT	2995
Db		4659	ATGGAATTAATATAGTGTCAATGAAGTATGAGTGGAAACCCCTTTCCTTACTTTTACCT	4718
Qy		2996	TCATTTCTTAGTATTAATTTTTTTTCTTACACCCCTGATCAAGCCACTAGTAGAGCACT	3055
Db		4719	TCATTTCTTAGT-----TTTTTTTTTCTTACACCCCTGATCAAGCCACTAGTAGAGCACT	4773
Qy		3056	ATCTGCTGCGAGCTATATATATGACTTTTACAGCAACAACATTTGCTGTGGCCTCTTTGG	3115
Db		4774	ATCTGCTGTGAGCTATATATGACTTTTACAGCAACAACATTTGCTGTGGCCTCTTTGG	4833
Qy		3116	GAAAGGAAACAGGATAGCAGGAGCTCAGGCTAGCAAGTCTGGACTCAACCTAAAGCCAG	3175
Db		4834	GAAAGGAAACAGGATAGCAGGAGCTCAGGCTAGCAAGTCT- GACTTTGCCCTAAAGCCAG	4892
Qy		3176	AGGCATGGTTGATAGCAGAGAAAGTGAGGCTCTTTCAAGTGGGTGTCTTAAATAATCA	3235
Db		4893	AGGCATGGTTGATAGCAGAGAAAGTGAGGCTCTTTCCGAAAGTGGGTGTCTTAAATAATCA	4952
Qy		3236	GAAACAGGAAGGCTCTGGTTGATGGAAATATACAGTAAAGATATCTACCCCTATCTCTTCT	3295
Db		4953	GAAACAGGAAGGCTCCGGTTGATGGAAATATACAGTAAAGATATCTACCCCTATCTC---CT	5009
Qy		3296	TCTATAGAAGCTAAACCGTCTCTCTCTTGTGTGTAGGCTGATAAACAACGCTTGTTTTC	3355
Db		5010	TCTATCGAACCTAAATCGTCTCTTTTCTTGTGTGTAGGCTGATAAACAACACTTGTTTTC	5069
Qy		3356	TTTTGAGTGTTCATGGCTTTGCAGATTTTCAGTGTCTCTGCCAGTTCCTTTGTAGAGGTTT	3415
Db		5070	TTTTGAGTGTTCATGGCTTTGTAGATTTTATAGTGTCTCTGCCAGTTCCTTTGTAGAGGTTT	5129
Qy		3416	GTTTACCTTTGACACCTGGCTTGGATGTAGCATGCCAAAGGCAACACTTCTGATGCCT	3475
Db		5130	GTTTACCTTTGACACCTGGGCTTGGATGTAGCATGCCAAAGGCAACACTTCTGATGCCT	5189
Qy		3476	GTGTAAAGGTTATATTACTTTTGTCTTTGGAAAGGTGAAGTGTGTGTGAGAAAG	3535
Db		5190	GTGTAAAGGTTATATTACTTTTGTCTTTGGAAAGGTGAAGTGTGTGTGAGAAAG	5249
Qy		3536	AACTCACAGGAGATGTATTTCTCTGTAGAAAC- TTTTTCCTCCCTTAAAGCCTATAA	3594
Db		5250	AACTCACAGGAGATGTGTCTCTGTAGAAACACTTTTTCCTCCCTTAAATGCTATAA	5309
Qy		3595	TCCACTTTCAGTCAACTTTGACTTTTATACATGCTGTCTACATGAAGAGTGTTAGGCC	3654
Db		5310	TCCACTTTCAGTCAACTTTGACTTTTATACATGCTGTCTACATGAAGAGTGTTAGGCC	5369
Qy		3655	CGCTCTCGTGTCTCTGGGAAAGCAACCAATAGGGGAGAAATGTTATGCCGAGAAATCTG	3714
Db		5370	CGCTCTCATGCTCTCTGGGAAAGCAACCAATAGGGGAGAAATGTTATGCTGAGAAATCTG	5429
Qy		3715	ACTGGCAGGAAACTGGGTTCAGAGCTCCCCAAAGACCACTACAGGTGTAAAGTAGGAACA	3774
Db		5430	ACCGCAGGAAACTGGGTTCAGAGCTCCCCGAAAGACCAACCAAGGTGTAAAGTAGGAACA	5489
Qy		3775	GTGAGGGTGGTTCATATATAGTAATGGAACAGAGGGGAGAGATAGCTACAAAGTT	3834
Db		5490	GTCCAGGGTGGGCTCATGTATATAGTAATGGAACAGAGCGAGGGAAGATAGCTACAAAGTT	5549
Qy		3835	TCATAGGCTCTTAAAGTATACAAATAGCTGGTGGCTTTCATAACAAAGGAAG	3894

Db		5550	TCATAGGCTCCGAGTCTTTAAAGATACAAATAGCTGCTTGGGCTTCATAACAAGGAAG	5609
Qy		3895	TCTGGGAGGCGACCAACCATTTAGAGGAGATCGAAAGGGGAAAAAC- -AATGTAGAGGA	3952
Db		5610	TCTGGGAGGCGACCAAG- ---TGAGAGGGAATGGAAGGGGAAAAAAGATGTAGAGGA	5666
Qy		3953	TTTGAAGAGCTACAAATCCTCCACGAGGAGATTTTCTTGGAGGAATCTAGAAACAAGGGT	4012
Db		5667	CTTGAACAGCTACAAATCCTCTACCAGACGATTTTCTTGGAAACAATCTAGAA- ---GGT	5722
Qy		4013	GGTGGATTAGGTGATCGCAGAGGACTTGTCTTGGCAATTTGAATCTGGGTTTTTGTCTC	4072
Db		5723	AGTGGATTAGGT- -GATTCAGAGGGGACTTGTCTTGGCAATTTGAATCTGGGTTTTTGTCTC	5781
Qy		4073	TCCATTTGAGGTTGAGAGCGTCAACCTTTTACCTCGATAGGAGGAGGAAGAGGGGT	4132
Db		5782	TCCATTTGAGGTTGAAAGCGTCAACCT- TTTTACCTCGAATGGAGGAGGAAGAGGGGT	5840
Qy		4133	GTTTGTACTCTTACTCGAGGTTTTTACTAGTTTACGCAATGGAAACAGACACTCGGGACCTC	4192
Db		5841	GTTTGTACTCTTACTCGAGGTTTTTACTAGTTTACGCAATGGAAACAGACACTCGGGACCTC	5900
Qy		4193	CTCTTGAAGAAAGGAGGAAAGGAAAGGAAAGGAAAGGAAAGGAAAGGAAAGGAAAGGAAAGG	4252
Db		5901	CTCTTGAC-----AAAAAATGGAAGCCTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTG	5950
Qy		4253	TTAAGAAAGCACAGGCGAGCTGGGCGATGGTGGCCCATGCCCTTTAATCCAGCAATTTGGGAG	4312
Db		5951	TTAAGAAAGCACCA-----	5963
Qy		4313	GCAGAGGCGAGTGACTTTCTAATTTCAAGGCCAGCTGGTCTACAAAGTGAGTTCCAGGA	4372
Db		5964	-----	5963
Qy		4373	CAGCCAGGCTATACAGAGAAACCCCTGTCTCGGAAAAAAGGAGAGAGAGAGAGAGAGAGAG	4432
Db		5964	-----	5963
Qy		4433	AAAGAAAG	4492
Db		5964	-----	5963
Qy		4493	GAA	4552
Db		5964	-----	5963
Qy		4553	GAGAAAGAA	4612
Db		5964	-----	5963
Qy		4613	AAAGAAAG	4672
Db		5964	-----GGCAAGCCCGACCATGGGT	5985
Qy		4673	CGTATGGGTCTTTGAGACAGAGCTTTTGAATTTGAGCGCTCATCAATAGTTGATCATGG	4732
Db		5986	TGAATGTGGGTCTTTGAGTCAAGGCTTTGAGTTGAGCACTCATCAATAGTTGATCATGG	6045
Qy		4733	TCAGGTGGAGGCTTACCTGTCAAGCCGAGCCCTGCTGGCTTGAACACTTCCAGG	4792
Db		6046	TCAGGTGGAGGCTTACCTGTCAAGCCGAGCCCTGCTGGCTTGAACACTTCCAGG	6105
Qy		4793	TCTCAGTATCACTTCTCTGCTTTAGCAGAGTTAGGAGTTGAGCAACCTTTTTCCTCA	4852
Db		6106	TCTCAGTATCACTTCTCTGCTTTAGCAGAGTTAGGAGTTGAGCAACCTTTTTCCTCA	6165
Qy		4853	CCCCCACTTAAATTTAATTTACAAAGCGAGTAAATTTGTGGGATACAGTGTGATTAAT	4912
Db		6166	CCCCCACTTAAATTTAATTTACAAAGCGTGTGTAATTTGTGGGATACAGTGTGATTAAT	6225
Qy		4913	GATCTATGTGTGCAATTTGTGCAAGGTTCAATAGGTTAGTCAATAGGCGCCATCAACAGCTT	4972
Db		6226	GATCTATGTGTGCAATTTGTGCAAGGTTCAATAGGTTAGTCAATAGGCGCCATCAACAGCTT	6285

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QY 4973 TATGGGTGTAAGTGAATATAGGTAGTGCCTGT-GTGTCTTAGGTGAGAAAG 5031
Db |||
QY 6286 TATGGGTGTAAGTGAATATAGGTAGTGCCTGTGTCTTAGGTGAGAAAG 6345
Db |||
QY 5032 CATGATTTTAAAGTCTTGGGCAATCATATATATCTCATGTGTTAAATGATTTGTTGA 5091
Db |||
QY 6346 CATGATTTTAAAGTCTTGGGCAATCATATATATCTCATGTGTTAAATGATTTGTTGA 6405
QY 5092 TTATCAATCTTTTAGAAGGCTGATCTTGTGTTTGGTCTCAGCAAGCAAAATGTCACC 5151
Db |||
QY 6406 TTATTAATCTTTTAGAAGGCTGATCTTGTGTTTGGTCTCAGCAAGCAAAATGTCACC 6465
QY 5152 AGCTCTTTTAACTAGTACCACTTTAGAAAATGCTACCCGTGCTCAAAATGGTTGTTGATT 5211
Db |||
QY 6466 AGCTCTTTTAACTAGTACCACTTTAGAAAATGCTACCCGTGCTCAAAATGGTTGTTGATT 6525
QY 5212 CTTATTTTCAATAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCT 5271
Db |||
QY 6526 CTTATTTTCAATAGCTTGGAGAGCGGAGAGATCAAAAGCGATCGGGAACTGGACCTGCT 6585
QY 5272 GTTTATCTCTGAGAAATGCTTGCCTGCTGAGCGAAGAAAGCTAGAAAACGAAGAACTG 5331
Db |||
QY 6586 GTTTATCTCTGAGAAATGCTTGCCTGCTGAGCGAAGAAAGCTAGAAAACGAAGAACTG 6645
QY 5332 CTCCTTCTCTCTTCTTAAAGAAACAATAAGATCCCTGAAATGGAATCTTTTACTAAAGGA 5391
Db |||
QY 6646 CTCCTTCTCTCTTCTTAAAGAAACAATAAGATCCCTGAAATGGAATCTTTTACTAAAGGA 6705
QY 5392 AAGTGAGAACTAAGCTCCACCATCATATAGAAATTTCAATGAAACCTGGCTCAGTTGA 5451
Db |||
QY 6706 AAGTGAGAACTAAGCTCCACCATCATATAGAAATTTCAATGAAACCTGGCTCAGTTGA 6765
QY 5452 AAGAGAAATAGTCTCAAGTTGTCATGAGACCAAGAGTAGACTTTGATAACCAAAAGAT 5511
Db |||
QY 6825 AAGAGAAATAGTCTCAAGTTGTCATGAGACCAAGAGTAGACTTTGATAACCAAAAGAT 6825
QY 5512 TCATTTGACAATATTTTATTTGTCATGATAATGCAACAGAAAAAGATGTAATCTTTTAAAAA 5571
Db |||
QY 6826 TCATTTGACAATATTTTATTTGTCATGATAATGCAACAGAAAAAGATGTAATCTTTTAAAAA 6885
QY 5572 TTGTTTGAAGAGGTTTACCTCTCATCTCTAGACCAAGAGTAGACTTTGATAACCAAAAGAT 5631
Db |||
QY 6886 TTGTTTGAAGAGGTTTACCTCTCATCTCTTGAAGAAAAAGCTTATGTAATCTTTTCAATTC 6945
QY 5632 CATAACCAATCTTTATATATGTAAGTTTATTTATATAGTATACATTTTATTTATGTC 5691
Db |||
QY 6946 CATATCCAAATTTTATATATGTAAGTTTATTTATATAGTATACATTTTATTTATGTC 7005
QY 5692 AGTTTATTAATATGGAATTTTATAGAAAAATTTATCTGATGTTGATATTTGATATATAA 5751
Db |||
QY 7006 AGTTTATTAATATGGAATTTTATAGAAAAATTTATCTGCTATTTGATATTT-AGTATAAG 7064
QY 5752 GCAAAATATTTATGATAATATATAGAAAAAAGATATCTTAGGCTTTTAAATAACACA 5811
Db |||
QY 7065 GCAAAATATTTATGATAATATATAGAAAAAAGATATCTTAGGCTTTTAAATAACACA 7124
QY 5812 TGAATATCAAAATCTTCTGCTGTAATTTTCTCCCTTTTAAATATCAAAATACCAATCA 5871
Db |||
QY 7125 TGGATATCAAAATCTTCTGCTGTAATTTTCTCCCTTTTAAATATCAAAATACCAATCA 7184
QY 5872 TCCTCATCATATACCAATCTTCTCATGCTTTCATGCTTGAAGTATTTCTGTTGTAAG 5931
Db |||
QY 7185 TCATCATCATATACCAATCTTCTCATGATTTTCATGCTTGAAGTATTTCTGTTGTAAG 7244
QY 5932 TTG 5935
Db |||
QY 7245 GTTG 7248
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RESULT 3  
US-11-177-987-26  
; Sequence 26, Application US/11177987

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; Publication No. US20050271619A1
; GENERAL INFORMATION:
; APPLICANT: Dumoutier, Laure
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac
; FILE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof
; FILE REFERENCE: LUD 5664
; CURRENT APPLICATION NUMBER: US/11/177,987
; CURRENT FILING DATE: 2005-07-07
; PRIOR APPLICATION NUMBER: US/09/626,617
; PRIOR FILING DATE: 2000-07-27
; PRIOR APPLICATION NUMBER: US09/419,568
; PRIOR FILING DATE: 1999-10-18
; PRIOR APPLICATION NUMBER: US09/354,243
; PRIOR FILING DATE: 1999-07-16
; PRIOR APPLICATION NUMBER: US09/178,973
; PRIOR FILING DATE: 1998-10-26
; NUMBER OF SEQ ID NOS: 43
; SEQ ID NO 26
; LENGTH: 4797
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
US-11-177-987-26

Query Match 11.0%; Score 650; DB 7; Length 4797;
Best Local Similarity 56.5%; Pred. No. 1e-133;
Matches 1863; Conservative 0; Mismatches 1285; Indels 152; Gaps 29;
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QY 356 CTCTCTCTCAGTTATCAACTTTTGACACTTGTGCGATCGGTGATGGCTGCTCTGCGAGAA 415
Db |||
QY 416 ATCTATGAGTTTTTCCCTTATGGGACTTTGGCGCGAGCTGCTGCTCTCTCATTTGCCCT 475
Db |||
QY 89 ATCTGAGCTCTTTCTTATGGGACCTTGGCCACAGCTGCTCTCTCTCTTTGGCCT 148
QY 476 GTGGGCGCCAGGAGGCAAAATGCGCTGCCCATCAACACCCGGTGCAAGCTTGAGGTGTCCAA 535
Db |||
QY 149 CTTGGTACAGGAGGAGCAGCTGCGCCCATCAGCTCCCATGTCAGAGCTTGACAAGTCCAA 208
QY 536 CTTCCAGAGCGGTATCATGCTCAACCGCACCTTTATGCTGGCCAGAGGAGGTACAGTGC 595
Db |||
QY 209 CTTCCAGAGCGGTATATCAACCAACCGCACCTTCATGCTGGCTTAAGGAGGTATACATCTC 268
QY 596 TCTCTTCTCTCCATACCGCTTGGCCATTTCTCTGAAGCACTTGCAAACTCTTTAGGGGC 655
Db |||
QY 269 AATCTGCTCTTTCTCGTTGGATCTACTTGGAAATCCAAATAGTTCTTTAAACTTTTCTTCA 328
QY 656 GCTTTATCTCCGAGGTCTCAGTACCTATGTTTTTCTGTCT- - - - -CTTTAGAG 703
Db |||
QY 329 GAGCATCTCTAAGAGCTTTTAGGAACCCAGCTTTATCCCTGAGGAGTAGATAAAATTTCTG 388
QY 704 ACTCTTTAAGGAGCTGATCTTTTCTATTTCTATTTCAAGGTCTCAGGACCATTTCTAT 763
Db |||
QY 389 TTTTTCAGAGACTCTTTGGGAATCTGGCTTTTTTTTTTTTCTTGAACCTCTTCTCTCCAT 448
QY 764 CTTGGGCTTCAGACACATATCTGAATTTTATCTACAGAGGCGCTTT- -AGAAAGCCA 821
Db |||
QY 449 TTTGGGCTTTATGATACATATGATGATTTTCCAAAGAGCGGCCATTCAGTAATCCAT 508
QY 822 CCCAGACTGCAATACTTTTCCATCTCTGTTGCTCTCTCTGAACTCATACTCTCTTGGC 881
Db |||
QY 509 CTGATGATTTTTTTTCTTATGCTCTGTCGATTTCTTAACTCATGCACACATCTG 568
QY 882 TACTC- - - - -CTGAGACCCACTCGGGAGACATACATCTCTAC 916
Db |||
QY 569 AATCTGCTTTTAGTCTTTTATGATGTTGCTCTGGGAGACGGGATGGGACATGCTAT 628
QY 917 TTACAGGCTTTTCTTCCATCTCTCTGTCACCCAGGACCTTAGGGTTTC- -TCTCTTTCAG 975
Db |||
QY 629 GTATAAATTTTTTTTCTATTTGCTCAATGTCCAGACCCCTTAGTCTTTTCTCTCTCCAG 688
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Qy	976	GCACGCTTGCAGATAACAACACAGACGTCGGCTCATACGGGAGAGAAATGTGTTCCAGGA	1033
Db	689	GCTAGCTTGGCTGATAAACAACACAGACGTTCTGCTCTCATTTGGGAGAGAAATGTGTTCCAGGA	748
Qy	1036	GTCACTGTAAAGTCCTCACTGCTGATGACAGGGC-----TAGCTGGGAGCT	1082
Db	749	GTCACTGTAAAGCTACAGTTGTGACGAACAGGGCGGTGTCGTCATGGGTACTTGGGGT	808
Qy	1083	GGTGGACCCCTCTGGGATAG----TCTGACGCTATGACCCCTGCTGCTTCTTGTCTACTCTGC	1138
Db	809	GCTGCTGATGATGTTTAAAGTCTTATCCCTTATGACCCCTTCTGTTTCCCTTCCACTGC	868
Qy	1139	AGGCTAAAGGATCAGTGTCTACCTGATGAAGCAGGTGCTCAACTTCAOCTTGGAGACATTC	1198
Db	869	AGATGAGTGAGCGCTGCTATCTGATGAAGCAGGTGCTGAACCTTCAOCTTGAAGAAGTGC	928
Qy	1199	TGCTCCCCAGTACAGACAGGTTCCGGCCCTACATGACAGGAGGTGTGTCCTTCTCTACCA	1258
Db	929	TGTTCCCTCAATCTGATAGGTTCCAGGCTTATATGCAGGAGGTGTGTCCTTCTCTGGCCA	988
Qy	1259	AACTCAGCAATCAGCTCAGCTCTGTGTAAAGTCTGGCTCTGGCTACCTATGCTCTCTCT	1318
Db	989	GGCTCAGCAACAGGCTAAGCAATGTGTAAAGTCTCAGCTCTCAGCCTATGCCACTTACCC	1048
Qy	1319	CTTCCTCTTCTATTTCCAGTAAGAACCCGAGGTCTGCCCCTCTCTCTTTCACAAGAGTGA	1378
Db	1049	CTCCTTCCCTCTTCCACAGAGACCCCTTACCCCAACTCTCTCTCTTCCCCTTACCCC	1108
Qy	1379	GGAGGCGCTCAGCACACACCATATAGGCCACTTGAATAGGTACAAAGGCTTTGGC	1438
Db	1109	TAACTACAGCAGGAAGAAGTGTCTTGGCAGCAGTGTATACAGGATCA-----TTTGGG	1161
Qy	1439	TTCAATTGAGTAATACCTTTGAGTTTGTATAGTTTAAAGCTTTATTTGTTTATCCATGGAA	1498
Db	1162	ATCATAGAGTATTTGCTTTTTCCTTTGACTGAGTCAOATCTTGAGTTTATAGTGGTGAATG	1221
Qy	1499	AGAAATCAACTCAAAATCTGTAGGATGAGAAAGATGTGGGAAACGAAAAAGCCCTAGAT	1558
Db	1222	GGGCTCGAACTTAAAGTGTACAGAAGCCGCAATTGGTTTGTCTTCGGAANAAGCAACTC	1281
Qy	1559	AGAGAAACAGATCTGCTGAGTACAGTACTTATAGGGGGGGGGGCGAGGGGGCGATATCCA	1618
Db	1282	AGGTTGCGTAA--GATGAGAAAGGTGTGTGGGAAACAATCTAGCTGTGTGGAATGGATCCA	1338
Qy	1619	CTGAGTCCAAGTACTGTGTGGGAGAGAAATCCACTGAGTACAGTACTCTGTGGGGAAGG	1678
Db	1339	TTGAGTCTAAAGTTGTGTAGGGGGGGGATGGCATGGAGAGAAATTAGAAGAGAAAGTGGG	1398
Qy	1679	AATGGCACAGACGAAAAAGTTCAAGGGGAAAGAGGAAGATGGAGAGCCCTCAATGTGGGGG	1738
Db	1399	AAATGGGNAAGCTTAAAGTCG-----GTGGTGGGTTCGCGACATGTTGCCCTGTGTA	1450
Qy	1739	TGTGAAAGGTCACTCTCTTTTCCATGTATGATGGAGAGTTAAGAAAAATCAGTGTGTGATTT	1798
Db	1451	TGTATGGGAAGCCACAAAATCGGAGGCGTGTGAACCTTGTATGTCGCGCTGAAACATTTGAAAC	1510
Qy	1799	TGATGTCTTACAGACCCCAACTATGCCAGACTGTGGGAGACCTGGGCAATTTAGGGA--AGG	1857
Db	1511	TATGAAAAAAGTTTGAGTGGAGTGGGCCCAAGTAAAAAGGCCCTTAGGACTTACTGGAAGGG	1570
Qy	1858	CGCGCTTTTCACACGAAAACTTTATGCTCATCTCTTGTGCTACACTCCCAOCTTTGAT	1917
Db	1571	GCTTAAATTTTACATGAGTGTTTTATGTACATTTCTTGTTCTAAGCATGCAATTTTCTG	1630
Qy	1918	CAGGTTAAAGCTCAGGTTTCGTTTCT-----ACGGTTCTTGTCTAC	1956
Db	1631	GAGATACCATTTAGGTTTATTTCTCTACAGAAATTTGCATAAACTACTCCGCTCTTTCCAC	1690
Qy	1957	TGGTGGAAACTTCAGTAGGATTTCCCAAGACGAGGACAGCTCTTCTGTAAAGGAGGAC	2016
Db	1691	AAATGCAAACCTCAGTAGGATTTCCCAAGATGAAGAGAGGTCTCTTGTAAAGGGAAGTGA	1750
Qy	2017	CTGGATTTTCACTGTCTTAGGAAACGAATAGCTCAGAGAAATCTTAGGTCAACGTTGAAATCT	2076

[illegible]

Db 2815 CTATAGTGAATGATACCATCATGTGGCCCTATTGTTGGTGAAGAAGACA--ACAATGGGAAGGC 2872  
QY 3141 TCAGGCTAGCAAGTCTCGACTCAACCTAAAGCCAGAGGCATGTTGATAGCAGAGAAAGT 3200  
Db 2873 TTAGACTAAACAATAGT-GACTCACCCCAAAACCGAGAGATGATTAGGACAGTGAAGT 2931  
QY 3201 GAGGCTCTTCAAGTGGGTGCTTAAAGTAATCAGAAAACAGGAAGGCTCTGTTGATGG 3260  
Db 2932 GAGGCTCTT-GCAAGCAGGTACAACTAAATTAATCTCAGAAAACATGAAGGCTCCAGTTGATGG 2990  
QY 3261 AATTATCAGTAAGATCTACCTTATCTCC-----TTCTCTATAGAGCTAAACCG 3313  
Db 2991 AATTTTCAGTAACAAGCTTAACCTTAATCCCCCTTTTCCCTCTGACTCTTTTAAAAA 3050  
QY 3314 TCTCTCCTCTTGTGTGTAGGCTGATAAACACGCTTGT--TTCTTTTGAAGTGTTCATGG 3371  
Db 3051 GCGTTTCTCTGAGCATCATTTAATGAGTGTGACTGTCTTCTCTTTGATAATGAGG 3110  
QY 3372 CTTTGCAGATTTTCAGTGTCTGCGAGTCTTGT--TAGAGGGTGTGTTACCTTGCACACC 3429  
Db 3111 CTTTGTAGTTTAAATTTGTGAAGCCAGTCTCTCTGTTATAGAACTATTATCTAGACATG 3170  
QY 3430 TGGGCTTGAATGTAGCATGCCAAAGGCACACACTCTGATGCCCTGTGTAAGAAGTTAT 3489  
Db 3171 GAGGCTGAATGTAGCATGCCACAGCAAGGATGCTTTACACATCTTCTGCTTAAAAAT 3230  
QY 3490 TATTCAATTACT-----TTGCTTTTGAAGGTGAAGTGTGTGAGAAAAGAACTCA 3541  
Db 3231 TACTGATTCACTTGTCTTGTCTTTAGAAAAGTGAAGTGTGAGAGAGAGAACTCA 3290

RESULT 4

US-11-177-987-9  
; Sequence 9, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LJD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 9  
; LENGTH: 1111  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-11-177-987-9

Query Match 10.1%; Score 602.4; DB 7; Length 1111;  
Best Local Similarity 99.8%; Pred. No. 1.5e-123;  
Matches 603; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
QY 5221 ATAGCTTGGAGAGCGGAGAGATCAAGCGATCGGGAACTGGACCTGCTGTTTATGTC 5280  
Db 508 AAAGCTTGGAGAGCGGAGAGATCAAGCGATCGGGAACTGGACCTGCTGTTTATGTC 567  
QY 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAAAGCTAGAAAACGAAGAACTGCTCTTCTCT 5340  
Db 568 TCTGAGAAATGCTTGGCTCTGAGCGAGAAAGCTAGAAAACGAAGAACTGCTCTTCTCT 627  
QY 5341 GCCTTCTTAAAGAACAAATAGATCCCTGATGGACTTTTCTTAAAGAAAGTGAGAA 5400

Db 628 GCCTTCTTAAAGAAACAATAAGATCCCTGAATGGAGCTTTTTTACTAAAGAAAGTGAGAA 687  
QY 5401 GCTAAAGCTCCACCATCATTTAGAGAGATTTTCAATGAAACCTGGCTCAGTTTGAAGAGAAAA 5460  
Db 688 GCTAAAGCTCCACCATCATTTAGAGAGATTTTCAATGAAACCTGGCTCAGTTTGAAGAGAAAA 747  
QY 5461 TAGTGTCAAGTTTGTCTCATGAGACGAGGTAGACTTTGATTAACCAAAAGATTTCAATTGACA 5520  
Db 748 TAGTGTCAAGTTTGTCTCATGAGACGAGGTAGACTTTGATTAACCAAAAGATTTCAATTGACA 807  
QY 5521 ATATTTTATTTGTCATTTGATAATGCAACAGAAAAAGTATGTACTTTTAAAAAATTTGTTTGA 5580  
Db 808 ATATTTTATTTGTCATTTGATAATGCAACAGAAAAAGTATGTACTTTTAAAAAATTTGTTTGA 867  
QY 5581 AGGAGGTTTACCTCTCATTTCTCTAGAGAAAAAGCCTATGTAACTTCATTTTCCATAACCAA 5640  
Db 868 AGGAGGTTTACCTCTCATTTCTCTAGAGAAAAAGCCTATGTAACTTCATTTTCCATAACCAA 927  
QY 5641 TACTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700  
Db 928 TACTTTATATATGTAAGTTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 987  
QY 5701 ATATGATATTTATTTATAGAAAAATTTATCTGATCTTGATATTTTCTGAGTATAAGCAAAATAAT 5760  
Db 988 ATATGATATTTATTTATAGAAAAATTTATCTGATCTTGATATTTTCTGAGTATAAGCAAAATAAT 1047  
QY 5761 ATTTATGATAATAACTATAGAAAAAGATATCTTAGGCTTTTAAATAACACATGAATATCA 5820  
Db 1048 ATTTATGATAATAACTATAGAAAAAGATATCTTAGGCTTTTAAATAACACATGAATATCA 1107  
QY 5821 TAAA 5824  
Db 1108 TAAA 1111

RESULT 5

US-11-177-987-7  
; Sequence 7, Application US/11177987  
; Publication No. US20050271619A1  
; GENERAL INFORMATION:  
; APPLICANT: Dumoutier, Laure  
; APPLICANT: Renaud, Jean-Christophe  
; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Fac  
; TITLE OF INVENTION: Interleukin-21, The Proteins Encoded, and Uses Thereof  
; FILE REFERENCE: LJD 5664  
; CURRENT APPLICATION NUMBER: US/11/177,987  
; CURRENT FILING DATE: 2005-07-07  
; PRIOR APPLICATION NUMBER: US/09/626,617  
; PRIOR FILING DATE: 2000-07-27  
; PRIOR APPLICATION NUMBER: US09/419,568  
; PRIOR FILING DATE: 1999-10-18  
; PRIOR APPLICATION NUMBER: US09/354,243  
; PRIOR FILING DATE: 1999-07-16  
; PRIOR APPLICATION NUMBER: US09/178,973  
; PRIOR FILING DATE: 1998-10-26  
; NUMBER OF SEQ ID NOS: 43  
; SEQ ID NO 7  
; LENGTH: 1119  
; TYPE: DNA  
; ORGANISM: Mus musculus  
; FEATURE:  
US-11-177-987-7

Query Match 9.4%; Score 555.2; DB 7; Length 1119;  
Best Local Similarity 96.0%; Pred. No. 4.1e-113;  
Matches 580; Conservative 0; Mismatches 23; Indels 1; Gaps 1;  
QY 5221 ATAGCTTGGAGAGCGGAGAGATCAAGCGATCGGGAACTGGACCTGCTGTTTATGTC 5280  
Db 510 AAAGCTTGGAGAGCGGAGAGATCAAGCGATTTGGGAACTGGACCTGCTGTTTATGTC 569  
QY 5281 TCTGAGAAATGCTTGGCTCTGAGCGAGAGAAAGCTAGAAAACGAAGAACTGCTCTTCTCT 5340

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Db 570 TCTGAGAAATGCTTGGCTCTGACGAGAGAGCTAGAAAAACGAAAGAACTGCTCCTTCCT 629
Qy 5341 GCCTTCTAAAGAAACAAATAGATCCCTGAAATGACATTTTTTCTAAAGGAAAGTGAGAA 5400
Db 630 GCCTTCTAAAGAAACAAATAGATCCCTGAAATGACATTTTTTCTAAAGGAAAGTGAGAA 689
Qy 5401 GCTAACGTCACCATCATATTAGAAGATTTCACATGAAACCTGGCTCAGTTGAAAAAGAAAA 5460
Db 690 GCTAACGTCACCATCATATTAGAAGATTTCACATGAAACCTGGCTCAGTTGAAAAAGAAAA 749
Qy 5461 TAGTGCAAGTTGTCCATGAGACAGAGGTAGACTTGATTAACACCAAGATTCATTGACA 5520
Db 750 TAGTGCAAGTTGTCCATGAGACAGAGGTAGACTTGATTAACACCAAGATTCATTGACA 809
Qy 5521 ATATTTTATGCTCATGATGATGCAACAGAAAAAGTATGACTTTAAAAAATGCTTTGAA 5580
Db 810 ATATTTTATGCTCATGATGATGCAACAGAAAAAGTATGACTTTAAAAAATGCTTTGAA 869
Qy 5581 AGGAGGTTACCTCTCATTTCTCTAGAGAAAGCCTATGTAACCTTCATTTCCATAACCAA 5640
Db 870 AGGAGGTTACCTCTCATTTCTCTAGAGAAAGCCTATGTAACCTTCATTTCCATAACCAA 929
Qy 5641 TACTTTATATGTAAGTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 5700
Db 930 TATTTTATATGTAAGTTATTTATTAAGTATACATTTTATTTATGTCAGTTTATTA 989
Qy 5701 ATATGGATTTATTTATAGAAAAATTTATCTGATGTTGATATTTGAGTATAAGCAAAATAT 5760
Db 990 ATATGGATTTATTTATAGAAACATTTCTGCTATGTTATTTT-AGTATAGGCAAAATAT 1048
Qy 5761 ATTTATGATATAACTATAGAAAAACAGATATCTTTAGGCTTTTAAATAACACATGATATCA 5820
Db 1049 ATTTATGACATAACTATGGAACAAGATATCTTTAGGCTTTTAAATAACACATGATATCA 1108
Qy 5821 TAAA 5824
Db 1109 TAAA 1112

RESULT 6
US-11-102-240-153
; Sequence 153, Application US/11102240
; Publication No. US20050260647A1
; GENERAL INFORMATION:
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Wood, William I.
; TITLE OF INVENTION: ANTIBODIES TO POLYPEPTIDES ENCODED BY A NUCLEIC ACID UNDEREXPRESS
; FILE REFERENCE: P3230R1C106C
; CURRENT APPLICATION NUMBER: US/11/102,240
; CURRENT FILING DATE: 2005-04-08
; PRIOR APPLICATION NUMBER: 10/063662
; PRIOR FILING DATE: 2002-05-07
; PRIOR APPLICATION NUMBER: 10/006867
; PRIOR FILING DATE: 2001-12-06
; PRIOR APPLICATION NUMBER: PCT/US00/23328
; PRIOR FILING DATE: 2000-08-24
; PRIOR APPLICATION NUMBER: 60/170262
; PRIOR FILING DATE: 199-12-09
; NUMBER OF SEQ ID NOS: 170
; SEQ ID NO 153
; LENGTH: 1152
; TYPE: DNA
; ORGANISM: Homo Sapiens
US-11-102-240-153

Query Match 3.6%; Score 215; DB 7; Length 1152;
Best Local Similarity 69.9%; Pred. No. 5.3e-38;
Matches 444; Conservative 0; Mismatches 155; Indels 36; Gaps 10;
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Qy 5221 ATAGCTTGGAGAGAGCGAGAGATCAAAGCGATCGGGGAACCTGGAACCTGCTGTTTATGTC 5280
Db 516 AAAAGCTTGGAGAGAGTGGAGAGATCAAAGCAATTTGGAGAACTGGAATTTGCTGTTTATGTC 575
Qy 5281 TCTGAGAAATGCTTGGCTCTGACGAGAGAGCTAGAAAAACGAAAGAACTGCTCCTTCCT 5340
Db 576 TCTGAGAAATGCTTGGCAATTTGACGAGACAAAGCTGAAAAATGAATTAACCAACCCCTTT 635
Qy 5341 GCCTTCTAAAAAGAAACAATAAGATCCCTGAATGACATTTTTTTT-----ACTAAAGGAAAGTG 5396
Db 636 CCCTGCTAGAAATAACAATTAGATGCCCAAGCGATTTTTTTTAAACCAAAAGGAAGATG 695
Qy 5397 AGAAGCTAAACGTCACCATCATTTAGAGATTTTCACTGAAACCTGGCTCAGTTGAAAGAG 5456
Db 696 GGAAGCCAAACTCCATCATGATGGTGGATTTCCAAATGAACCCCTGCGTTAGTTACAAAG 755
Qy 5457 AAAATAGTGTCAA--GTTGTCCATGAGACCCAG--AGGTAGACTTGTATTAACCAACCAAGATTC 5513
Db 756 GAAACCAATGCCACTTTTGTATTATAGACCAAGGAGGTAGACTTTCCTAAGCATAGATATTT 815
Qy 5514 ATTGACAATATTTTATTGTCTATTGATA---ATGCAACAGAAAAAGTATGTACTTTTAAAA 5569
Db 816 ATTGATAACATTTTCAATTTGTAACCTGGTGTCTATACACAGAAAAACAATTTATTTTAAAT 875
Qy 5570 AATTGTTT-----GAAAGGAGGTACCTCTCATCTCCTCTAGAGAAAA---CCCTATG 5619
Db 876 AATTGCTTTTTCATAAAAAAGATTACTTTTCCATCTCTTAGGGGAAAAAACCCCTAAA 935
Qy 5620 TAACTTCA--TTTCCATAACCATACTTTATATATGTAAGTTTATTTATTAAGTATA- 5676
Db 936 TAGCTTCATGTTTCCATAAATCAGTACTTATATTTATAAATGATATTTATTTATTATATAA 995
Qy 5677 -----CATTTTATTATGTCAGTTTATAATATGATTTTATTTATAGAAAAATTTATCTGA 5731
Db 996 GACTGCAATTTTATTATATCATTTTATATATGATTTTATTTATAGAAACATCATTCGA 1055
Qy 5732 TGTGATATTTTGAGTATAAGCAAAATA---ATATTTATGATAATAACTATAG-----AAAC 5784
Db 1056 TATTGCTACTTGGTGAAGGCTAATATTGATATTTTATGACATAAATATTAGAGCTATAA 1115
Qy 5785 AAGATATCTTAGGCTTTAAATAACACATGATATC 5819
Db 1116 CATGTTTATTGACCTCAATAAACACATTTGGATATC 1150

RESULT 7
US-11-121-086-83
; Sequence 83, Application US/11121086
; Publication No. US20050266459A1
; GENERAL INFORMATION:
; APPLICANT: POULSEN, TIM S.
; APPLICANT: NIELSEN, KIRSTEN V.
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES
; FILE REFERENCE: 09138.6000-00000
; CURRENT APPLICATION NUMBER: US/11/121.086
; CURRENT FILING DATE: 2005-05-04
; PRIOR APPLICATION NUMBER: 60/567,570
; PRIOR FILING DATE: 2004-05-04
; NUMBER OF SEQ ID NOS: 107
; SOFTWARE: Patent in version 3.3
; SEQ ID NO 83
; LENGTH: 18745
; TYPE: DNA
; ORGANISM: Homo sapiens
US-11-121-086-83

Query Match 2.7%; Score 159.4; DB 7; Length 187745;
Best Local Similarity 64.1%; Pred. No. 1.4e-24;
Matches 257; Conservative 0; Mismatches 141; Indels 3; Gaps 1;
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Qy 4256 AGAAGACACAGGCACTGGGCGATGGTGGCCCATCCCTTTATTCACGATTTGGGAGGCA 4315
Db 41385 AAAATAAAAAATTAGCCAGGCACTGGTGGCATATGCTGTGGTCTCAGCTACTTGGGAGGCT 41444
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Qy	4316	GAGCGAGTGACATTTCTAAATTC	CAAGCGCAGCCTGCTCTACAAAGTGAGTTC	CCAGGACAG	4375
Db	41445	AAGCGAGGAGGATCGCTT	AGGCG---TTCAGTGAAGCCATGTT	CATGACACTGCATTC	41501
Qy	4376	CCAGGGGCTATACAGAGAAACCC	TGTCGCGGAAAAA	AAAAAAGAGAAAGAAAAAGAA	4435
Db	41502	CCTGGGTGACAGAGACCC	TATCAGAAAGGAGAAAGAGAAAGGGGAGG	AGAGAGAGGGG	41561
Qy	4436	AGAAGAGAAAGAGAGAGGAGAG	GAGAGGAGAGAGAGAGAGAGAGAGAGAGAG	4495	
Db	41562	ACGAGAGGAGAGCGGAGGAGAG	GAGAGGGGAGCGGAGAGAGAGAGAGAGAGAGAG	41621	
Qy	4496	AGGAGGAGAGAGAGAGAGAGAG	AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG	4555	
Db	41622	AAGGGGAGAGGGGAGAGAGAC	CGGGAGGAGAGCGGGAGGAGAGCGGGAGGAGAGGGGAGAG	41681	
Qy	4556	AAGAGAGAGAGAGAGAGAGAG	AAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG	4615	
Db	41682	AGGGGAGGAGGGGAGGAGAG	GGGGAGGAGAGGGGAGGAGAGGGGAGGAGGGGAGGAGGGG	41741	
Qy	4616	AGAAAAAGAAAAAAGCAAGCA	AGCAAGCACTGGCAAA	4656	
Db	41742	AGAAAGGAAAGAGGAGAAAG	AGAAAGAGAGAAAGCAAGCAAACTG	CACAA 41782	

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RESULT 8
US-10-333-025-24
; Sequence 24, Application US/10933025
; Publication No. US20050265987A1
; GENERAL INFORMATION:
; APPLICANT: ROSEN, STEVEN
; APPLICANT: HEMMERICH, STEFAN
; APPLICANT: TOMITA, MEGUMI
; TITLE OF INVENTION: Sulfotransferases and methods of use

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	Query Match	2.4%	Score 145;	DB 6;	Length 171936;
	Best Local Similarity	69.8%;	Pred. No. 2e-21;		
	Matches 196;	Conservative 0;	Mismatches 85;	Indels 0;	Gaps 0;
Qy	4255	AGAAAGCA CAGCAGCTGGGCATGCTGGCCCATGCCTTTAATCCAGCATTTGGAGGC	4314		
Db	71997	AAAGAAAGAAAGAAAGTCAAGCGCTGGTGGCGCATGCTTTAGTCCAGCACATTTGGAGGC	72056		
Qy	4315	AGAGCGAGGTGACTTCTTAAATTCAAGGCGAGCGCTGGTCTCAAAAGTGAAGTTCAGGACA	4374		
Db	72057	AGAGGACGGCAGATTTCTGAGTTCAGAGGCGAGCGCTGGTCTACAGAGTGAAGTTCAGGACA	72116		

Qy	4375	GCACGGCTTATACAGAGAAAACCTGTCTCGGGAAAAA	4434	GAAGAAAGAAAGAAAGAAAGAA
Db	72117	GCACGGCTCATATAGAGAAAACCTGTCTCTAAACCC	72176	AAAAAGAAAGAAAGAAAGAAAGAGAGAGAGAG
Qy	4435	AAGAAGAAAGAGGAGAGAGAGAGAGAGAGAGAGAG	4494	AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG
Db	72177	AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG	72236	AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG

[illegible]

Query Match	2.4%;	Score 142;	DB 6;	Length 26209;
Best Local Similarity	68.8%;	Pred. No. 1.1e-20;		
Matches 231;	Conservative	0;	Mismatches 90;	Indels 15;
Gaps 2;				

Qy 4368 CAGGACAGCCAGGGCTATACAGAGAAACCTGTCTCGGGAATAAAAAAGAGAAAG 4427

[illegible][illegible]

Qy 4548 GAGAAGAGAAGAGAAGAGAAGAGAAG 4583  
|||  
Db 47517 GAGAAGAGAAGAGAAGAGAAGAGAAG 47552

```

RESULT 10
US-11-176-253-1
; Sequence 1, Application US/11176253
; Publication No. US20050262579A1
; GENERAL INFORMATION:
; APPLICANT: Yamanouchi Pharmaceutical Co., Ltd.
; TITLE OF INVENTION: Novel clock gene promoter
; FILE REFERENCE: Q75308

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[illegible]





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